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GenCore version 5.1.9 Copyright (c) 1993 - 2006 Biocceleration Ltd.

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1598 segs, 35149 residues Searched:

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SUMMARIES

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	Reverse transcript Reverse transcript	transcrip	rse transcri	e transcrip	Reverse transcript Reverse transcript	e transcri	Reverse transcript Reverse transcript	se transcri	transcri	Reverse transcript Reverse transcript	everse transcri	se transcri	YMOTPhic	ase Oligo	ranscrip	transcri		transcr	transcrip	DNA anscr	e transcri	e transcri	Reverse transcript Reverse transcript	se transcri	everse transcri	กัด	a)	Reverse transcript	Reverse transcript Reverse transcript	Reverse transcript	Reverse transcript Reverse transcript	Synthetic oligonuc	Human dendritic ce	Human oligonucleot	Human oligonucleot	Human oligonucleot	Human oligonucleot	AA463249-derived o	Human stanniocalci AIO34360-derived o	AI041482-derived o	AA679352-derived o	Human giucocortico Chimeric phosphoro	Human mPGES-1 chim	Human mitoNEET-spe	keverse transcript Reverse transcript	everse tran	everse trar	Reverse transcript Reverse transcript	everse tran
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Reverse transcript Reverse transcript Reverse transcript Reverse transcript Reverse transcript	Human oligonucleot Human glucocortico	Human glucocortico Chimeric phosphoro	Chimeric phosphoro	Reverse transcript	Reverse transcript Reverse transcript	Reverse transcript	keverse transcript Reverse transcript	transcri	transcri	transcri	transcri	transcri	נו נ	transcrip	transcri	Reverse transcript	Reverse transcript	Reverse transcript	Reverse transcript Reverse transcript	Reverse transcript	Reverse transcript	ψq	ַ פַ	0 :	4 4	. ⋖	E .	Rat DNA microarray	A Bi	A microarra	CpG immunostimulat	ICAM-1 bind	2',5'-Inned tetra Reverse transcript	_	- +		transcri	trans	Reverse transcript Reverse transcript	trans	transcri	Reverse transcript	transcri	transcri	trans	everse transcri	everse transcri	Reverse transcript	everse transcri
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Anti-proliferative Antisense oligonuc Antisense oligonuc Antisense oligonuc Antisense oligonuc 5' mRNA DNA prepar Hepatitis C virus Hepatitis C virus Human Fas and Fasi Human Pas and Fasi Human ICAMI SiRNA	HUMAN 1 LCM1 B1RNA TYT18Apad PS27-20- CardiovasCular dis HUMAN 0 1900nucleot AA25835-derived o HUMAN glucocortico Chimeric phosphoro Chimeric phosphoro HUMAN IL-10 PCR pr Acetyl-Coenzyme A- HUMAN EZH2 SINA 10- HUMAN EZH2 SINA 10- HUMAN HER2 (EGFR2) HUMAN HERA INFOSINE P COW PRION PROTECEIN HUMAN GLUCOSE-6-Dh	Glucose-6-phosphat XIAP targeting siR Muman IL-4R siRNA Human IL-4R cansc	an NOGO receg an chondrocyt detecting pr detecting pr rogen recept rogen recept rogen recept rogen recept an MDZ7 scann an MDZ7 scann	CpG immunostimulat CpG immunostimulat CpG immunostimulat CpG immunostimulat CpG immunostimulat IL-10 expression a IC-10 express
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Example 1; SEQ ID NO 31969; 47pp; English.

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splice variant; transcriptome; oligonucleotide library; ss.
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ADC49403
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ADT64925
AAQ30448
AAQ30447
                                          AAV54172
AAV54171
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02-MAY-2001; 2001US-0287724P.
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(COMP-) COMPUGEN INC.
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New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and developmental-specific genes.

Faigler S;

Mintz L,

Mintz E,

Shoshan A, Wasserman A,

WPI; 2002-257383/30.

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The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptom comprises messenger RNAs transcribed from multiple comprises messenger RNAs transcribed from multiple comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a proposical sample, in expression profiling studies, in qualitatively or quantitatively characteristing the corresponding transcriptom, and in cetecting RNA transcripts and splice variants of human or animal transcripts and splice variants of human or animal content and pathological state, and so allowing the detection of tissue and pathology-specific genes such as those genes only expressed in competing to tissue under a specific pathological condition; to detect condition; to devect conditions to developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular cissue condition; to ABNESSESS perpresent oligonucleotide sequences from disorder. ABNESSES perpresent oligonucleotide sequences from crats, humans and mice, which are used in the exemplification of the principle of the printed specification, but was obtained in electronic format diagrantly, from units of the circuit, in the condition of the manner of the print of the circuit, in the condition of th
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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02-MAY-2001; 2001US-0287724P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shoshan A, Wasserman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (COMP-) COMPUGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200210449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN59222;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Shoshan A,
                                      ABN59220;
                          Query Match
                            Matches
                                   RESULT 3
                                    ABN59220
ઠે
                                셤
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New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and developmental-Example 1; SEQ ID NO 6003; 47pp; English Shoshan A, Wasserman A, (COMP-) COMPUGEN INC. WPI; 2002-257383/30. specific genes. Matches ABN33255 8 The present invention describes oligomuclectide libraries for detecting messenger RNAs that populate a (sub-)transcribtome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligomuclectides, each capable of hybridising selectively to a set of messenger RNAs transcribtion unit of the genome, consequence on or more messenger RNA splice variants. The coligomuclectide libraries are useful for detecting manAs from a coligomuclectide libraries are useful for detecting manAs from a coligomuclectide libraries are useful for detecting manAs from a coligomuclectide libraries are useful for detecting manAs from a coligomuclectide libraries may also be used as specialised minical state, and splice variants of human or animal colibraries to detect transcripts of a sub-transcriptome under a particular bloogical or pathological state, and so allowing the detection of tissue colification or pathological state, and so allowing the detection of tissue specific tissue under a specific pathological condition; to detect can specific genes; and to detect RNA transcripts and splice condition; to developmental specific genes; and to detect RNA transcripts and splice developmental specific genes; and to detect RNA transcripts and splice developmental specific genes; and to detect RNA transcripts and splice cates, humans and mice, which are used in the exemplification of the printed specification, but was obtained in electronic format chirach prom with are used in the sequence data for this patent did not form directly from with an in the min of the principal produced to the printed specification, but which are used in the sequences from directly from with an in the produced and performed the principal produced to the produced t 2564 TICTCCTGAGCTAGGAAGAGICTACCCGACATAAGTCGAGGGACTTTATGTTTTTGAGGC 2623 New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of a genome, useful for detecting tissue-, pathology-, and developmental-1 TCTCCTGAGCTAGGAAGAGTCTACCGACATAAGTCGAGGACTTTATGTTTTGAGGC 60 0; Gaps Human spliced transcript detection oligonucleotide SEQ ID NO:31968 Human; mouse; rat; splice transcript; detection; RNA transcript; splice variant; transcriptome; oligonucleotide library; ss. directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences Score 60; DB 1; Length 60; Pred. No. 2.4; 0; Indels Mintz E, Mintz L, Faigler S; Sequence 60 BP; 14 A; 12 C; 16 G; 18 T; 0 U; 0 Other; Similarity 100.0%; Pred. No. 2.4 50; Conservative 0; Mismatches Example 1; SEQ ID NO 31970; 47pp; English. Example 1; SEQ ID NO 31968; 47pp; English. 20-JUL-2001; 2001WO-IB001903. 28-JUL-2000; 2000US-0221607P. 02-MAY-2001; 2001US-0287724P. ABNS9220 standard; DNA; 60 Wasserman A, (COMP-) COMPUGEN INC WPI; 2002-257383/30. WO200210449-A2. specific genes. Local Sim Homo sapiens 15-JUL-2002 07-FEB-2002

The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptom comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The coligonucleotide libraries are useful for detecting mRNAs from a coligonucleotide libraries are useful for detecting mRNAs from a coligonucleotide libraries are useful for detecting mRNA transcripts and splice variants of human or animal constitutions. The libraries may also be used as specialised minical libraries to detect transcripts of a sub-transcriptome under a particular colibraries to detect transcripts of a sub-transcriptom under a particular coliological or pathological state, and so allowing the detection of tissue specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient suffering from a particular colisconder. ABN73753 to ABN75589 represent oligonucleotide sequences from cates, humans and mice, which are used in the exemplification of the print of the printed specifician of the printed specification, but was obtained in ole form directly from with a specification, but was obtained in olectronic format directly. 1051 ö 9 992 GCAAAACGAATTCTAGAGCTTGACCAGTTTAAGGGGCAGCAGGGACAAAAACGTTTCCAA GCAAAACGAATTCTAGAGCTTGACCAGTTTAAGGGGCAGCAGGGACAAAAAAGGTTTCCAA 0; Gaps Human spliced transcript detection oligonucleotide SEQ ID NO:6003. Human, mouse, rat, splice transcript, detection, RNA transcript, splice variant, transcriptome, oligonucleotide library, ss. directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences Score 60; DB 1; Length 60; 0; Indels Sequence 60 BP; 22 A; 12 C; 15 G; 11 T; 0 U; 0 Other; 100.0%; Pred. No. 2.4; ive 0; Mismatches ABN33255 standard; DNA; 60 BP 28-JUL-2000; 2000US-0221607P. 02-MAY-2001; 2001US-0287724P. 20-JUL-2001; 2001WO-IB001903 (first entry) Local Similarity 100. nes 60; Conservative WO200210449-A2 Homo sapiens. 15-JUL-2002 07-FEB-2002. ABN33255; Query Match

Faigler S;

Mintz L,

Mintz E,

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messenger RNAs that populate a (sub-) transcriptome, where the (sub-
transcriptom units that populate a genome. The library comprises several
coligonucleotides, each capable of Pybridising selectively to a set of
coligonucleotides, each capable of Pybridising selectively to a set of
coligonucleotides, each capable of Pybridising selectively to a set of
messenger RNAs transcribed from a given transcription unit of the genome,
thich encodes one or more messenger RNA splice variants. The
coligonucleotide libraries are useful for detecting mRNAs from a
coligonucleotide libraries are useful for detecting mRNAs from a
coligonucleotide libraries are useful for detecting mRNAs from a
constitutively characterising the corresponding transcriptom, and in
certing RNA transcripts and splice variants of human or animal
cranscriptomes. The libraries may also be used as specialised mini
clibraries to detect transcripts of a sub-transcriptom under a particular
chlological or pathological state, and so allowing the detection of tissue
chlological or pathological state, and so allowing the detection of tissue
collogical or pathological state, and so allowing the detection of tissue
collogical or pathological state, and so allowing the detection of tissue
collogical or pathological state, and to detect RNA transcriptom of a gracial defect of developmental specific genes; and to detect RNA transcripts and splice
collogical cranscriptome of a patient suffering from a particular
clibrarians of a transcriptome of a patient suffering from a particular
clibrarians and mice, which are used in the exemplification of the
consent invention. N.B. The sequence date for this patent did not form
clibraries better the prince of t
present invention describes oligonucleotide libraries for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Sequence 60 BP; 14 A; 13 C; 20 G; 13 T; 0 U; 0 Other;

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                                                                      2119 GAACCTGGAGGCCTTGGCTTTGATTGATGGAGCCGGAACAAGCAGTGGACCTGACATT 2178
                                                                                      1 GAACCTGGAGGCCTTGGCTTTGGATTTGATGGAGCCGGAACAAGCAGTGGACCTGACATT 60
                                            Gaps
                                            ö
             DB 1; Length 60;
2.4;
                                         0; Indels
Match 2.2%; Score 60; ub ... Local Similarity 100.0%; Pred. No. 2.4
               Query Match
                                            Matches
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Human spliced transcript detection oligonucleotide SEQ ID NO:31796. Human, mouse, rat, splice transcript, detection, RNA transcript, splice variant, transcriptome, oligonucleotide library, ss. ABN59048 standard; DNA; 60 BP (first entry) WO200210449-A2. Homo sapiens 15-JUL-2002 07-FEB-2002 ABN59048; RESULT 5 ABN59048 

20-JUL-2001; 2001WO-IB001903.

28-JUL-2000; 2000US-0221607P. 02-MAY-2001; 2001US-0287724P.

(COMP-) COMPUGEN INC

New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of genome, useful for detecting tissue-, pathology-, and developmental-Faigler S; Mintz L, щ Mintz Wasserman A, WPI; 2002-257383/30. Shoshan A,

Example 1; SEQ ID NO 31796; 47pp; English.

specific genes.

The present invention describes oligonucleotide libraries for detecting

CC messenger RNAs that populate a (sub-)transcriptome, where the (subltranscriptome comprises messenger RNAs transcribed from multiple
cc transcription units that populate a genome. The library comprises several
coligomucleotides, each capable of bybridising selectively to a set of
messenger RNAs transcribed from a given transcription unit of the genome,
which encodes one or more messenger RNA splice variants. The
coligomucleotide libraries are useful for detecting mRNAs from a
coligomucleotide libraries are useful for detecting mRNAs from a
coligomucleotide libraries are useful for detecting mRNAs from a
coligomucleotide libraries may also be used as specialised mini
cdetecting RNA transcripts and splice variants of human or animal
cranscriptomes. The libraries may also be used as specialised mini
clibraries to detect transcripts of a sub-transcriptom under a particular
cbiological or pathological state, and so allowing the detection of tissue
clibraries to detect transcripts of a sub-transcriptom and splice
closelopmental specific genes; and to detect RNA transcripts and splice
covariants of a transcriptome of a patient suffering from a particular
clibraries and mice, which are used in the exemplification of the
crats, humans and mice, which are used in the exemplification of the
coligoral pathology. Proceeding in the exemplification of the
coligomucleotide sequences from
crats, humans and mice, which was obtained in electronic format
coliforal. 2564 TICICCIGAGCIAGGAAGAGICIACCCGACATAAGICGAGGGACTITATGITTIGAGGC 2623 9 Gaps ö directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences Match 2.2%; Score 60; DB 1; Length 60; Local Similarity 100.0%; Pred. No. 2.4; es 60; Conservative 0; Mismatches 0; Indels Sequence 60 BP; 14 A; 12 C; 16 G; 18 T; 0 U; 0 Other; Query Match Matches à 셤

1 TTCTCCTGAGCTAGGAAGAGTCTACCCGACATAAGTCGAGGACTTTATGTTTTTGAGGC

BP ADC22315 standard; DNA; 54 (first entry) 18-DEC-2003 ADC22315; RESULT 6 ADC22315

recombinant fusion protein; fusion protein; binding; detection; Nuclear localisation signal nucleotide sequence SEQ ID NO:164. subcellular compartment localisation; gene; ds. localisation domain; binding domain;

01-AUG-2002; 2002WO-US024572. 01-AUG-2001; 2001US-0309395P. 13-DEC-2001; 2001US-0341589P. WO2003012068-A2. Homo sapiens. 13-FEB-2003. 

Chen Y; Premkumar DR, WPI; 2003-248174/24. Bright G,

P-PSDB; ADC22314.

(CELL-) CELLOMICS INC.

New recombinant fusion protein comprising detection and first localization domains and a binding domain for the molecule of interest, useful for detecting binding of a molecule of interest.

Claim 20; SEQ ID NO 164; 101pp; English

The present invention describes a recombinant fusion protein (I) for

detecting binding of a molecule of interest. (I) comprises: (a) a detection domain; (b) a first localisation domain; and (c) a binding comain for the molecule of interest. The detection domain the first localisation domain and the binding domain for the molecule of interest constituting the recombinant fusion protein for detecting binding of a molecule of interest are operably linked. The binding domain for the molecule of interest is separated from the first localisation domain by 0 -20 amino acid residues. The first localisation domain by 0 c-20 amino acid residues. The first localisation domain by 0 c-20 amino acid residues. The first localisation domain for the molecule of interest both do not occur in a single non-recombinant for the molecule of interest. Also described:

(1) a recombinant nucleic acid encoding the recombinant fusion protein;

(2) a recombinant expression vector comprising the nucleic acid control sequences operably linked to the recombinant nucleic acid molecule; (3) a genetically engineered host cell transfected with the recombinant control sequences operably linked to the recombinant mucleic acid molecule; (3) a genetically engineered host cell transfected with the recombinant control interest; and (5) a method for identifying compounds that alter the interest; and (5) a method for identifying compounds that alter the binding of the molecule of interest. The recombinant control is useful for detecting binding of a molecule of interest. The recombinant controls and enables the molecule of construct two or more ofineering the construct who or more ofineering and enables the molecule of construct two or more ofineering the construct when an enables in the molecule of interest in the molecule of interest. proteins and enables the monitoring of biochemical events in live, intact or fixed cells. The present sequence is used in the exemplification of the present invention. 

Sequence 54 BP; 25 A; 9 C; 13 G; 7 T; 0 U; 0 Other;

Gaps ; 2.0%; Score 54; DB 1; Length 54; 100.0%; Pred. No. 5.9; tive 0; Mismatches 0; Indels 54; Conservative Similarity Query Match Local Matches ò

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2270 AAAGTTACCAAGAGAAACACGATAATGAAGGTTCTGGAAGCAAAAGGCCCAAG 2323 54 AAAGTTACCAAGAAAAACACGATAATGAAGGTTCTGGAAGCAAAAGGCCCAAG

g

BP ABZ06784 standard; DNA; 50 (first entry) 09-JAN-2003 ABZ06784; ABZ06784/c 

Human leukocyte gene expression profiling probe SEQ ID NO 6775.

17; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;

Homo

WO200257414-A2.

25-JUL-2002

22-OCT-2001; 2001WO-US047856.

20-OCT-2000; 2000US-0241994P. 08-JUN-2001; 2001US-0296764P.

(BIOC-) BIOCARDIA INC

Prentice J, Phillips J; K, Matcuk G, Altman P, Pr Quertermous T, Johnson F; nuth J, Fry K, Matcuk G, Woodward R, Quertermous T Wohlgemuth J, ľ, S

WPI; 2002-636525/68.

New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.

Claim 1; Page 547; Opp; English

The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ0815) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual. The diseases include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis or cytomegalovirus infection

Sequence 50 BP; 10 A; 8 C; 15 G; 17 T; 0 U; 0 Other;

1542 ; 1493 GCTCTCAAGCCTCCTCCAATAAAGCTCTATCGGGAAACAAATGAACCAGT 1.8%; Score 50; DB 1; Length 50; 00.0%; Pred. No. 11; 0; Indels Similarity 100.0%; Pred. No. ... 50; Conservative Query Match Best Local Matches ð

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Human leukocyte gene expression profiling probe SEQ ID NO 6385. -GCTCTCAAGCCTCCAATAAAGCTCTATCGGGAAACAAATGAACCAGT BP ABZ06394 standard; DNA; 50 (first entry) 09-JAN-2003 ABZ06394; 20 ABZ06394 a 

17, leukocyte, gene expression profiling, allograft rejection; atherosclerosis, congestive heart failure, systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, cytomegalovirus, infection, probe;

Homo sapiens.

WO200257414-A2.

25-JUL-2002.

22-OCT-2001; 2001WO-US047856.

20-OCT-2000; 2000US-0241994P. 08-JUN-2001; 2001US-0296764P.

(BIOC-) BIOCARDIA INC.

Phillips J; Prentice J, Ly N, Woodward R, Quertermous T, Johnson F; Altman P, Matcuk G, Fry K, Wohlgemuth J,

WPI; 2002-636525/68.

New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.

Claim 1; Page 536; Opp; English.

The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligomuclectides (ABZ00010-ABZ00152) each having 50 base pairs (Dp). The system is useful for leukocyte expression profiling. It is particularly useful for for leukocyte expression profiling (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual. The diseases include cardiac allograft

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                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence polymorphism analysis; human identity; human relatedness;
single nucleotide polymorphism; SNP; genetic disease; cytostatic;
immunosuppressive, antiinflammatory; neuroprotective; antimicrobial;
fatty acid metabolism; glycolysis; amino acid metabolism;
paternity analysis; forensic; autoimmune disease; cancer; nervous system;
rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis or cytomegalovirus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide and polypeptide including one or more single nucleotide polymorphisms, useful for diagnosing and treating conditions associated with the presence of sequence polymorphism in humans and
                                                                                                                                            Gaps
                                                                                                                                                                             1493 GCTCTCAAGCCTCCTCCAATAAAGCTCTATCGGGAAACAAATGAACCAGT 1542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /standard_name= "Single nucleotide polymorphism"
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0
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                                                                                                                                                                                                   GCTCTCAAGCCTCCTCCAATAAAGCTCTATCGGGAAACAAATGAACCAGT
                                                                                                                                                                                                                                                                                                                                                                                                               Human single nucleotide polymorphism (SNP) region Seq ID143.
                                                                                                         DB 1; Length 50;
                                                                                                                                            Indels
                                                                        Seguence 50 BP; 17 A; 15 C; 8 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection; pathogenic microorganism; human; ds
                                                                                            1.0%; Scor.
100.0%; Pred. No. 11.
... 0; Mismatches
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                                                                                                                                                                                                                                                                                                         BP.
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99US-00443199
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                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                            50; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-399731/34.
                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 variation
                                                                                                                                                                                                                                                                                                                                            ADC17041;
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                                                                                                                                            Matches
                                                                                                                                                                                                                                                                        RESULT 9
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metabolism, glycolysis, and amino acid metabolism may also be developed. The compounds may be useful for treating a subject suffering from or at risk for a pathology associated with the presence of a sequence polymorphism. SNP detection is also useful in paternity analysis and forensic application. Polymorphisms may contribute to the phenotype of an organism and phenotypic traits include genetic diseases such as autoimmune diseases, cancer, diseases of the nervous system and infection by pathogenic microorganisms. The present sequence is the sequence surrounding and including a human SNP of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence polymorphism analysis; human identity; human relatedness; single nucleotide polymorphism; SNP; genetic disease; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; fatty acid metabolism; glycolysis; amino acid metabolism; paternity analysis; forensic; autoimmune disease; cancer; nervous system; infection; pathogenic microorganism; human; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to novel isolated nucleotide sequences which comprise 217 defined polymorphic sequences. Sequence polymorphism-based analysis of nucleic acid sequences can augment or replace previously known methods for determining the identity and relatedness of individuals. Single nucleotide polymorphisms (SNPs) tend to occur with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel polynucleotide and polypeptide including one or more single nucleotide polymorphisms, useful for diagnosing and treating conditions associated with the presence of sequence polymorphism in humans and
                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                1618 CTATGGGAGTCGTCAGATTATACTGGAGAAAGAGGAAACAGAAGAGCTAAA 1668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /standard_name= "Single nucleotide polymorphism"
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0
                                                                                                                                                                                                                                                                                                                  51
                                                                                                                                                                                                                                                                                                    Human single nucleotide polymorphism (SNP) region Seq ID142.
                                                                                                                                                                                                              51;
                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                           Score 49.4; DB 1; Length
Pred. No. 12;
                                                                                                                                                                            Sequence 51 BP; 22 A; 6 C; 14 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 142; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers replace (26, C)
                                                                                                                                                                                                           1.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                        ADC17040 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                  50; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leach MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-399731/34.
                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200029622-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
variation
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC17040;
                                                                                                                                                                                                                Query Match
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great frequency throughout the genome and may be located close to loci of interest. Such variations can cause or be closely linked to pathological conditions (genetic diseases). Hence the SNPs of the invention may be useful in the development of compounds with cytostatic, may be activities. Regulators of metabolic pathways such as fatty acid metabolism, glycolysis, and amino acid metabolism may also be developed. The compounds may be useful for treating a subject suffering from or at risk for a pathology associated with the presence of a sequence polymorphism. SNP detection is also useful in paternity analysis and forensic application. Polymorphisms may contribute to the phenotype of an organism and phenotypic traits include genetic diseases such as autoimmune diseases, cancer, diseases of the nervous system and infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T7; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Phillips J;
                                                                                                                                                                                                                                                                     by pathogenic microorganisms. The present sequence is the sequence surrounding and including a human SNP of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                               1373 GAAGACCTGTTGCGGAAGGTTCGCGCCAAGGAGCCAGGAAGCGAGCACTC 1423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAGACCTGTTGCGGAAGGTTCGCGACAAGGAGCCAGGAAGCGAGCACTC 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human leukocyte gene expression profiling probe SEQ ID NO 82.
                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
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                                                                                                                                                                                                                                                                                                                             Sequence 51 BP; 15 A; 12 C; 18 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                           ed. No. 12;
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                      Score 49.4;
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                                                                                                                                                                                                                                                                                                                                                                    1.8%;
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08-JUN-2001; 2001US-0296764P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                       50; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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predicting disease complications in an individual or monitoring response to treatment in an individual. The diseases include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis or cytomegalovirus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, to treatment in an individual. The disease include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17, leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nitoring (the rate of) progression of a disease, e.g. atherosclerosis congestive heart failure, comprises diagnostic oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Phillips J;
                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rheumatoid arthritis, osteoarthritis or cytomegalovirus infection
                                                                                                                                                                                                              2629 GITGCCATGGTGATGGTGAGCCCTCCCACTTTGCTGTTCTTTAC 2678
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Human leukocyte gene expression profiling probe SEQ ID NO 2130.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New system for leukocyte expression profiling, diagnosing
                                                                                                                                                                               1; Indels
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                                                                                                                                            DB 1; Length
                                                                                                         Sequence 50 BP; 6 A; 14 C; 11 G; 19 T; 0 U; 0 Other;
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                                                                                                                              Score 48.4; DE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Altman P,
                                                                                                                                                                               0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nuth J, Fry K, Matcuk G, A
Woodward R, Quertermous T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 394; Opp; English.
                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-OCT-2000; 2000US-0241994P.
08-JUN-2001; 2001US-0296764P.
                                                                                                                                            1.8%;
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                                                                                                                                                           Similarity
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                                                                                                                                                                               49;
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ABZ00089;

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The present invention relates to a method (M1) for detecting and/or isolating a nucleic acid having a homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) comprises treating a sample containing nucleic acid compounds with an locked nucleic acid acid compounds with an locked nucleic acid acid having the homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) is useful for detecting and isolating nucleic acids released from a lysed complex biological mixture comprising nucleic acids. The present sequence is a LNA capture probe, used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting and/or isolating nucleic acid molecule having homopolymeric sequence or repetitive element or conserved nucleotide sequence involves treating sample containing nucleic acid compounds with locked nucleic
                                                                                                                                                                                                                                                /*tag= a /mod_base= OTHER /mod_base= OTHER /note= "5' AQ2, where AQ is anthraquinone" | 16. .35 /*tag= b /*tag= b
                                                                                                                                  Detection; isolation; locked nucleic acid; LNA; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 1.3%; Score 35; DB 1; Length 35; Local Similarity 100.0%; Pred. No. 95; loservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "Optionally LNA nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 35 BP; 0 A; 0 C; 0 G; 35 T; 0 U; 0 Other;
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                                                                                                                                                                                                            Location/Qualifiers
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                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid oligonucleotide.
                                                                                             probe #1
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                                                         03-JUN-2004
                                                                                               LNA capture
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                                                                                                                                                                           Synthetic
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                ADL33738;
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                                                                                                                                                                                                                                                                                                                                  17; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
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    Gaps
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                                                                             20
                                                                                                                                                                                                                                                                                             Human leukocyte gene expression profiling probe SEQ ID NO 80.
                                        GTGATGGTGTAGCCCTCCCACTTTGCTGTTCTTTACTTTACTGCCTGAAT
                                                                               GTTGCCATGGTGATGGTGTAGCCCTCCCACTTTGCTGTTCTTTAC
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bred. No. 14;
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    Mismatches
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Woodward R, Quertermous T,
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llarity 98.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-OCT-2000; 2000US-0241994P.
08-JUN-2001; 2001US-0296764P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-OCT-2001; 2001WO-US047856
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                                                                                                                                                                             ABZ00089 standard; DNA; 50
                                                                                                                                                                                                                                                           (first entry)
    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOC-) BIOCARDIA INC.
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nes 49; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200257414-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wohlgemuth J,
                                                                                                                                                                                                                                                                                                                                                                                                                                        ношо варіепв
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    49;
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Gaps

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ò a ADL33738/c

RESULT 14

Query Match

Matches

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GENOVOXX GMBH
                                                                                                                                                                                                                          CHERKASOV D.
                                                                                                                                                                                                                                                                                                         WPI; 2006-040183/05
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                                                                                                                   DE102004025746-A1
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                                                   modified base
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                                                                                                                                             15-DEC-2005
             Synthetic.
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                                                                                                                                                                                                                                                      (GENO-)
                                                                                                                                                                                                                          (CHER/)
                                                                                                                                                                                                                                        (HENN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 17
                                        Key
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윱
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                                                                                                                                                               'note= "OTHER = Linked to unidentified chemical group in he absence of bases 32-35" 12.\ .35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to novel conjugates comprising 1-100 nucleotide or nucleoside molecules coupled to a label through water-soluble polymer linkers. The conjugates of the invention may be useful for modifying nucleic acid chains via coupling reactions catalyzed by DNA polymerase, RNA polymerase or terminal transferase enzymes or by phosphoramidite coupling, e.g. for labeling nucleic acids in arrays bound to a solid phase. The current sequence is that of the Cy3-labeled polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New conjugates useful for modifying nucleic acid chains comprise nucleotide or nucleoside molecules coupled to a label through water-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel solid phase-related oligonucleotide Oligo dT35-Cy3 #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 35; 95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 35 BP; 0 A; 0 C; 0 G; 35 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743
                                                                                                                 note= "OTHER = Labeled with Cy3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          35 АААААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.3%; Score ...,
100.0%; Pred. No. 95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modification PCR primer 2 of the invention
                                                                                                                                                                                                              /*tag= c
/note= "Optionally absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 34B; Page 123; 212pp; German.
                                                            Location/Qualifiers
                                                                                     *tag= a
/mod_base= OTHER
                                                                                                                                                        mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                05-NOV-2003; 2003DE-01051636
05-DEC-2003; 2003DE-01056837
                                                                                                                                                                                                                                                                                                     05-NOV-2004; 2004WO-EP012556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEE86826 standard; DNA; 35
                                                                                                                                         *tag= b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 1.3
Best Local Similarity 100.
Matches 35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         soluble polymer linkers.
                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                             Cherkasov D, Hennig C;
                                                                                                                                                                                                                                                                                                                                                                     (GENO-) GENOVOXX GMBH.
(CHER/) CHERKASOV D.
(HENN/) HENNIG C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-372185/38.
         PCR; primer; ss.
                                                                                                                                                                                             misc difference
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                                                                         modified base
                                                                                                                            modified base
                                  Unidentified
                                                                                                                                                                                                                                                                            19-MAY-2005
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This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged and respected contact of the solid phase with high concentrations of labeled reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; dna detection; DNA sequencing; DNA amplification; oligo dT35-Cy3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.3%; Score 35; DB 1; Length 35; 100.0%; Pred. No. 95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 35 BP; 0 A; 0 C; 0 G; 35 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743
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/mod_base= OTHER
/note= "OTHER= Cy3"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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/mod_base= 5'-Cy3
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ADN35261;
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                                                                                                                                                                                                              washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process that the process able to determine many sequences in their claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dT35-Cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                  This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus; RNA virus; replication; viral infection; PCR primer;
                                                                                                                                     Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                          1.3%; Score 35; DB 1; Length 35; 100.0%; Pred. No. 95;
                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ohmori H;
                                                                                                                                                                                                                                                                                                                        Sequence 35 BP; 0 A; 0 C; 0 G; 35 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus PCR primer CAC-T35 SEQ ID NO:19
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                                                                                                Baeuml E;
                                                                                                                                                                 Example 8; Page 90; 141pp; German
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                  26-MAY-2004; 2004DE-10025696
                                      26-MAY-2004; 2004DE-10025696
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                Cherkasov D, Hennig C,
                                                                  HENNIG C.
GENOVOXX GMBH.
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                                                         CHERKASOV D.
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                                                                                                                   WPI; 2006-185820/20
                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-1999;
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 23-FEB-2006
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                                                         CHER/)
                                                                             (GENO-)
                                                                    (HENN/)
                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                     The present invention describes a vector comprising a cDNA encoding an RNA virus gene, constructed to ensure the exact and homogeneous transcription of both terminals of the RNA virus gene. Also described is a method for screening drugs for inhibiting the replication of RNA virus by using the RNA viral infection model animal, particularly one with hepatitis C viral infection. The vector is useful in clarifying the mechanism of RNA viral replication, onset of RNA viral infection, and developing remedies and therapeutics for RNA viral infection, and particularly of a hepatitis C virus. The present sequence represents a PCR primer which is used in the exemplification of the present invention
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Vectors expressing full-length gene of RNA viruses, useful in clarifying mechanisms of RNA viral replication, infection, and developing remedies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           secondary-ion mass spectrometry, gene analysis; disease diagnosis; species identification; ss; probe.
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0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 38;
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Pred. No.
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                                                                                                                                                   Example 2; Page 20; 46pp; Japanese.
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28-JUN-2002; 2002JP-00191391.
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Best Local Similarity
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                                                                             and therapeutics
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Tue Nov

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WPI; 2000-338599/29
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                          (CHIR ) CHIRON
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modified_base
                                                                                                                 Unidentified
                                                                                                                                                                                                                               12-MAY-1995;
                                                                                                                                                                                                                                                                    28-SEP-1992;
                                                                                                                                                                                                                                                                                       28-SEP-1993;
                                                                                                                                                      US6057093-A.
                                                                                                                                                                                          02-MAY-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35;
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                                                                                                                                                                                                                                                                                                                                                              Han JH,
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       NAME OF THE PROPERTY OF THE PR
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                         for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV; thiolation; reverse transcriptase; primer; inhibition; homooligomer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       region (d(T)12) to promote the binding of the remaining portion of the E-thiolated oligonucleotride (MdV 28) to a homopurine site of the viral genome via triple-helix formations. The oligo is used to in the treatment of HIV. See also AAQ22624-27 and AAQ25017-Q25032
applicable in gene analysis, disease diagnosis and species identification. The present sequence represents a probe sequence used hybridization tests.
                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The oligomer comprises a non-thiolated (binding) homooligonucleotide
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                                                                                                               Length 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.3%; Score 35; DB 1; Length 40;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40 BP; 0 A; 0 C; 0 G; 12 T; 28 U; 0 Other;
                                                                         Sequence 39 BP; 0 A; 0 C; 0 G; 38 T; 1 U; 0 Other;
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                                                                                                             DB 1;
. le+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide specific for HIV proviral DNA.
                                                                                                                                                                                                                1.3%; Score 35; DB
100.0%; Pred. No. 1e+
:ive 0; Mismatches
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ID AAQ25031 standard; DNA; 40
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ID AAA39649 standard; DNA; 40
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                                                                                                                             Local Similarity 100.
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les 35; Conserv
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                                                                                                           Query Match
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This invention describes a novel method for enhancing translation of a coding region which involves making an RNA molecule, comprising the coding region operably linked to a 5' untranslated region (UTR) to comprising a sequence fully homologous to pestivirus homology box IV, and then translating it so that the translation of the coding region is enhanced. The method is useful for enhancing or controlling the translation of HCV nucleic acid, which allows stronger immune responses, where blocking or decreasing translation of viral nucleic acid may decrease the pathology of viral infection. This sequence represents a primer which is used in the construction of hybrid CAT RNA's described in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Enhancing translation of coding region of hepatitis C virus involves making RNA molecule comprising the coding region and 5' untranslated region comprising a sequence fully homologous to pestivirus homology
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                                                                Control element, translation enhancer, pestivirus homology box IV, immune response; viral infection; primer; ss.
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100.0%; Pred. No. 1e+02;
iive 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "5' HS-(CH2)6-O-PO2-O-T"
Primer used in construction of hybrid CAT RNA.
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93US-00128583.
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DNA microarray; gene analygis; disease diagnosis; species identification;
probe; ss.
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100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; SEQ ID NO 1; 59pp; Japanese.
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                                                                                                                                                                                                                                                                                             Aiba T,
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28-JUN-2002; 2002JP-00191390.
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                                                                                                                                                                                                                                                                                             rakase H, Okamoto T,
                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-083158/08.
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hes 35; Conserv
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ADH76858/c
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The invention relates to a DNA microarray for detecting a target nucleic acid molecule in a specimen, comprising a nucleic acid probe that has a target endernee substantially complementary to the base sequence of the target nucleic acid molecule immobilised onto a substrate. The microarrays are applicable in gene analysis, disease diagnosis and species identification. With such reliable microarrays, the detection of target nucleic acid molecules can be conveniently and quickly achieved, with high accuracy and improved reproducibility and quantitation even in a high-throughput system. In the microarray with a matrix shape, it is possible to shown images of distribution of formation density of a nucleic acid probe-dot system. Various internal and external-strandard synches were prepared biologically, by enzyme digestion or cleavage, and synthetically, for immobilisation onto a glass substrate e.g. after screen-printing at defined density distribution in a pattern. The current
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence represents a probe related to the invention.
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                                                                   Example 10; SEQ ID NO 9; 52pp; Japanese.
and species identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a probe support containing a nucleic probe region having nucleic probes fixed to it. Also described is a probe support analysis method. The probe support can be used for the TOF-SIMS analysis of nucleic acid probes by forming a phosphorus-containing region usable as a standard on a substrate, nucleic probes located as a matrix on a nucleic acid chip substrate can be accurately and quantitatively analysed. The support enables accurate analysis. The present sequence represents a probe which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe support contains nucleic probe region having nucleic probes fixed therein and enables accurate analysis of probes using TOP-SIMS.
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                                                                                                                                                                                                                                           nucleic acid base sequence detection; target nucleic acid; annealing; hybrid; dideoxy nucleotide triphosphate; deoxynucleotide triphosphate;
                             Gaps
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                                                                                                                                                                                                                     Nucleic acid sequence detection-related oligonucleotide SeqIDl.
  DB 1; Length 40;
1e+02;
                           0; Indels
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/note= "OTHER= Bound to O-PO2-(CH2)6-SH"
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               100.0%; Pred. No. letive 0; Mismatches
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   Score 35;
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/mod_base= OTHER
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                                                                                                                                            ADK70561 standard; DNA; 40
                                                                                                                                                                                              (first entry)
Query Match
Best Local Similarity 100.0
Matches 35, Conservative
                                                                                                                                                                                                                                                                        mass spectrometry; ss.
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                                                                                                                                                                                                                                                                                                                                     modified base
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                                                                                                                                                                                                                                                                                                Unidentified
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                                                                                                                                                                    ADK70561;
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                                                                                                                                   ADK70561
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This invention relates to a novel method of detecting a nucleic acid base

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sequence, which comprises fixing on board a partial structure cut by light within the 5' side of primer, adding a target nucleic acid as a template, performing annualing to form a hybrid, extending the hybrid using four sorts of dideoxy nucleotide triphosphate, deoxynucleotide triphosphate, removing the template, irradiating with light and analysing the sequence of extension part based molecular weight obtained by mass spectrometry. The invention is useful for determining the base sequence of a number of nucleic acids effectively and efficiently in a short time.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to a novel analytical method to target nucleic acid molecules in a sample. Specifically, it refers to a DNA probe carrier that has two or more types of probe that are complementary to the base sequence of the target RNA molecule such that the hybrid structure can subsequently be identified by secondary ion mass spectroscopy time of flight. The present invention describes a method that overcomes the problems associated with using radioactive isotopes or fluorescence to identify the nucleic acid molecules of interest. Accordingly, this method provides a means for the acquisition of accurate gene information. This oligonucleotide sequence is a target RNA sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis of target nucleic acid e.g., RNA or DNA, by making sample react with probe carrier having probes complementary to target and detecting the hybrid by time-of-flight type secondary ion mass spectrometry.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA probe carrier; secondary ion mass spectroscopy time of flight; ss; nucelic acid detection.
                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/mod_base= OTHER
/note= "OTHER= HS-(CH2)6-O-PO2-O, optionally absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA sequence target of novel nucleic acid detection method SeqID 1.
                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                           Length 40;
                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                     Sequence 40 BP; 0 A; 0 C; 0 G; 40 T; 0 U; 0 Other;
                                                                                                                                                                                                                      Score 35; DB 1; Ler
Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                    40 ААААААААААААААААААААААААААААААА
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100.0%; Pred. No. --
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                 exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK67293 standard; RNA; 40
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                                                                                                                                                                                                                                                                            35; Conservative
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                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of target nucleic acid e.g., RNA or DNA, by making sample react with probe carrier having probes complementary to target and detecting the hybrid by time-of-flight type secondary ion mass spectrometry.
                                                                                                                                                                                                                                                                                    carrier; secondary ion mass spectroscopy time of flight; ss;
                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/mod_base= OTHER
/note= "OTHER= HS-(CH2)6-0-PO2-0, optionally absent"
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  DB 1; her.,
0. 1e+02;
0; Indels
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Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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    1.3%; bcc.
100.0%; Pred. No. 100.0%; 0; Mismatches
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Query Match
Best Local Similarity 100.0
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Best Local Similarity 100.0
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                                                                                                                                                                                                                                                                                 probe
                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                              ADK67292;
                                                                                                                                      RESULT 26
                                                                                                                                                                                                                                                                                   DNA
                                                                                                                                                      ADK67292
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RESULT 27 ADJ71299/c

Sequence 40 BP; 0 A; 0 C; 0 G; 0 T; 40 U; 0 Other;

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This invention relates to a novel method for developing nucleic acid chip substrates. Specifically, it refers to a chip that comprises a nucleic acid probe and related nucleic acid substances arranged in a matrix form on the chip substrate surface. The present invention describes observing binding to the nucleic acid probe by an environment-control type scanning electron microscope and determining the quality of the dot shape produced. This oligonucleotide sequence is a DNA probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid chip substrate, comprising nucleic acid probe and related nucleic acid substances arranged in matrix form on substrate surface.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid chip substrate; probe; matrix;
environment-control type scanning electron microscope; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 35; DB 1; Length 40;
Pred. No. 1e+02;
0; Mismatches 0; Indels
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/note= "OTHER= HS-(CH2)6-O-PO2-O label"
                                                                                                        /mod_base= OTHER
/note= "OTHER= HS-(CH2)6-O-PO2-O label"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 40 BP; 0 A; 0 C; 0 G; 40 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 1; 17pp; Japanese.
                                                   Location/Qualifiers
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Best Local Similarity 100.
Matches 35; Conservative
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modified_base
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               Synthetic
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                                                     Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method of analysing a substance on a substrate using mass spectrometry (MALDI-TOF MS). The method involves using a structure including a partial structure to be disconnected by light to fix the substance on the substrate, irradiating the substance partial structure to be disconnected by light, and analysing the mass spectrum of the substance which is brought in an unfixed state by disconnecting the partial structure by the irradiation of light. The method can be used to acquire data from a bio-chip. The present sequence is a probe used to demonstrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Acquiring mass of immobilized substance, using mass spectrometry, by fixing substance to substrate using partial structure to be disconnected by light in bonded part, analyzing mass spectrum of substance after disconnecting structure.
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0
                                                                                                              Method of analysing substance using mass spectrometry probe #1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.3%; Score 35; DB 1; Length 40;
100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
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/note= "modified by HS-(CH2)6-0-PO-O"
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                                                                                                                                                  ss; probe; mass spectrometry; MALDI-TOF MS; light
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Labelled DNA oligonucleotide probe SeqID 1.
                                                                                                                                                                                                                             Location/Qualifiers
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ADJ71299 standard; DNA; 40 BP
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/mod_base= (
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                                                                           (first entry)
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Best Local Similarity 100.
Matches 35, Conservative
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modified_base
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                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Testing surface state of organic device in manufacturing organic device has organic film containing organic compound having sample holding base
                                                                                                                                    comprising nucleic acid probe and related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                               substances arranged in matrix form on substrate surface
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100.0%; Pred. No. 1e+02;
:ive 0; Mismatches 0
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                                                                                                                             Nucleic acid chip substrate,
nucleic acid substances arrar
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(CANO ) CANON KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a probe immobilised substrate and a method for its manufacture. The probes can be located using a surface analysis method e.g. scanning electron microscopy, atomic force microscopy, and time-of-flight secondary ion mass spectrometry (TOP-SIMS) method. The invention also discloses a method for analysing the probe array, and a method for detecting the probe and target material. The invention is useful for probe immobilised substrates such as DNA probes such as single strand DNA probes, single strand RNA probes, single strand peptide nucleic acid (PNA) probes, protein probes for genome analysis and peptide nucleic acid (PNA) probes, protein probes for genome analysis and determination of therapeutic strategies, using electron microscopy, photoelectron spectroscopy, atomic force microscopy, and TOP-SIMS. The
                   s analysed repeatedly in a high
sequence is a labelled DNA probe
arranged in matrix form on the substrate. In the manufacturing process, the laminated film of the biochip can be analysed repeatedly in a high throughput manner. This oligonucleotide sequence is a labelled DNA probused in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe immobilised substrate; surface analysis method; scanning electron microscopy; time-of-flight secondary ion mass spectrometry; TOF-SIMS; probe array; genome analysis; gene expression analysis; cancer; hereditary disease;
                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Probe immobilized substrate e.g. for DNA probe, has colored metal or metal compound provided at prescribed spots for locating probe spots.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Probe immobilised substrate method associated DNA probe #26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infectious disease; photoelectron spectroscopy; probe; ss.
                                                                                                                       1.3%; Score 35; DB 1; Length 40;
100.0%; Pred. No. 1e+02;
                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/mod_base= OTHER
/note= "Modified by thiol (SH) group"
                                                                                      Sequence 40 BP; 0 A; 0 C; 0 G; 40 T; 0 U; 0 Other;
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0
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                                                                                                                                                                                                                 100.0%; Pred. no.
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                                                                                                                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUN-2002; 2002JP-00189836.
28-JUN-2002; 2002JP-00190009.
                                                                                                                                                                                                                                                                                               ADM16960/c
ID ADM16960 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-338612/31.
                                                                                                                                   Local Similarity
es 35; Conserv
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                      03-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nakamura K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                   ADM16960;
                                                                                                                     Query Match
                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                RESULT 31
 8888888
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Matches
                                                                     RESULT 33
ADN35263/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to detecting a probe located and/or a target capable of binding specifically to the probe on a substrate comprises the preparation of a substrate with the probe and/or the target for specific binding to the probe on its surface, and measurement of the substrate surface by time- of-flight secondary-ion mass spectrometry with labeling. The method is for analyzing probe supports or nucleic acids on nucleic acid chips with detection and quantitation of probe conditions and hybrid of probe with target nucleic acid, which is applicable in gene analysis, disease diagnosis and species identification. The present sequence represents a probe sequence used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
invention reduces the time for searching the positions of probe array spots, hence reliable images are obtained quickly, thereby enabling efficient and accurate analysis, at low cost. The present sequence represents a DNA probe used in the exemplification of the present
                                                                                                                                                                                    Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          secondary-ion mass spectrometry; gene analysis; disease diagnosis; species identification; 88; probe.
                                                                                                                                                                                  ;
0
                                                                                                                                  DB 1; Lens.
3. 16+02;
0; Indels
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                                                                                                             Sequence 40 BP; 0 A; 0 C; 0 G; 40 T; 0 U; 0 Other;
                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743
                                                                                                                                                                                                                                                      40 hahahahahahahahahahahahahahah 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe sequence used for hybridization tests #1.
                                                                                                                                             1.3%; Score 35; DB 1
100.0%; Pred. No. 1e+C
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; SEQ ID NO 1; 68pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hashimoto H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JUN-2002; 2002JP-00190010.
28-JUN-2002; 2002JP-00191391.
28-JUN-2002; 2002JP-00191414.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUN-2003; 2003WO-JP008104
                                                                                                                                                                                                                                                                                                                                                   ADN35256 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Окатото Т, Такаве Н,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-203385/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JAN-2004
                                                                             invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                      ADN35256;
                                                                                                                                                                                                                                                                                                                 RESULT 32
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The present invention relates to detecting a probe located and/or a target capable of binding specifically to the probe on a substrate comprises the preparation of a substrate with the probe and/or the target for specific binding to the probe located on its surface, and measurement of the substrate surface by time- of-flight secondary-ion mass spectrometry with labeling. The method is for analyzing probe supports or nucleic acids on nucleic acid chips with detection and quantitation of probe conditions and hybrid of probe with target nucleic acid, which is applicable in gene analysis, disease diagnosis and species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                     secondary-ion mass spectrometry; gene analysis; disease diagnosis; species identification; ss; probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 40 BP; 0 A; 0 C; 0 G; 35 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 ААААААААААААААААААААААААААААААААААА
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                                                                                                                                                                                                                                                                                                          Probe sequence used for hybridization tests #5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hashimoto H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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28-JUN-2002; 2002JP-00191391.
28-JUN-2002; 2002JP-00191414.
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                                                                                                                                                                                                                                                           01-JUL-2004 (first entry)
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                                                                                                                                                         ADN35263 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Okamoto I, Takase H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-203385/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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Gaps

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1.3%; Score 35; DB 1; llarity 100.0%; Pred. No. 1e+02; Conservative 0; Mismatches 0

Local Similarity hes 35; Conserv

Matches

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Query Match

Length 40; 0; Indels

Synthetic

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The present invention relates to detecting a probe located and/or a target capable of binding specifically to the probe on a substrate comprises the preparation of a substrate with the probe and/or the target for specific binding to the probe located on its surface, and measurement of the substrate surface by time- of-flight secondary-ion mass spectrometry with labeling. The method is for analyzing probe supports or nucleic acids on nucleic acids chips with detection and quantitation of probe conditions and hybrid of probe with target nucleic acid, which is applicable in gene analysis, disease diagnosis and species identification. The present sequence represents a probe sequence used for hybridization tests.
                                                                                                                                                                                     Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.3%; Score 35; DB 1; Length 40; 100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 40 BP; 0 A; 0 C; 0 G; 35 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                       Example; SEQ ID NO 9; 68pp; Japanese
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                                                                                                       Hashimoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADN35265 standard; DNA; 40 BP.
28-JUN-2002; 2002JP-00191391 28-JUN-2002; 2002JP-00191414
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28-JUN-2002; 2002JP-00191391.
28-JUN-2002; 2002JP-00191414.
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species identification; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               35; Conservative
                                                                                                     Okamoto T, Takase H,
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                                                                                                                                               WPI; 2004-203385/19.
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                                                           (CANO ) CANON KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                 secondary-ion mass spectrometry; gene analysis; disease diagnosis;
species identification; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          spectrometry; gene analysis; disease diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.3%; Score 35; DB 1; Length 40; 100.0%; Pred. No. 1e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                            Hashimoto
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                                                                                                                                                                                                                                                                       28-JUN-2002; 2002JP-00191391.
28-JUN-2002; 2002JP-00191414.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADN35264 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          secondary-ion mass spectron species identification; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for hybridization tests.
                                                                                                                                                                                                                                                                                                                                                                            Takase H,
                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-203385/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
es 35; Conserv
                                                                                                                                                                                                                                                                                                                                  (CANO ) CANON KK.
                                                                                                                     WO2004003532-A1.
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                                                                                                                                                               08-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                      Okamoto T,
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Synthetic.

ADN35264;

ADN35264/c

Query Match

Best Loca Matches

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Gaps

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Example; SEQ ID NO 10; 68pp; Japanese,

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The present invention relates to detecting a probe located and/or a target capable of binding specifically to the probe on a substrate comprises the preparation of a substrate with the probe and/or the target for specific binding to the probe located on its surface, and measurement of the substrate surface by time- of-flight secondary-ion mass spectrometry with labeling. The method is for analyzing probe supports or nucleic acids on nucleic acid chips with detection and quantitation of probe conditions and hybrid of probe with target uncleic acid, which is applicable in gene analysis, disease diagnosis and species identification. The present sequence represents a target sequence used for hybridization tests.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Probe for measuring device of intermolecular interaction of organic
molecule, comprises probe portion coated with electroconductive film and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified nucleic acid used in intermolecular interaction measurement #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a probe comprising a probe portion (3) coated with an electroconductive film (4) and an organic film (5) by chemical bonds through a sulfur atom. Further disclosed is a method for manufacture of the probe, and a method for measuring intermolecular interactions. The method of the invention is useful for creating a measuring device e.g. scanning probe microscope (SPM) and atomic force microscope (AFM) used for measuring an intermolecular interaction in samples, such as organic molecules e.g. deoxyribonucleic acid (DNA).
                                                                                                                                                                                                                                                                                        1.3%; Score 35; DB 1; Length 40;
100.0%; Pred. No. 1e+02;
ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
/note= "HS-(CH2)6-0-PO2-0-Adenine"
                                                                                                                                                                                                                                                       Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                    2709 дадададададададададададададададада 2743
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Intermolecular interaction; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAR-2004; 2004JP-00099903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAR-2004; 2004JP-00099903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AED51679 standard; DNA; 40
                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0
Matches 35, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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modified_base
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organic film
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEDS1679
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The invention relates to a probe comprising a probe portion (3) coated with an electroconductive film (4) and an organic film (5) by chemical bonds through a sulfur atom. Purther disclosed is a method for menthod for measuring intermolecular interactions. The method of the invention is useful for creating a measuring device e.g. scanning probe microscope (SPM) and atomic force microscope (APM) used for measuring an intermolecular interaction in samples, such as organic molecules e.g. deoxyribonucleic acid (DNA). Probes of the invention enable stable measurement of organic molecules fixed on substrate by a simple structure. The current sequence represents a nucleic acid sequence used in an example of the invention.
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Probes of the invention enable stable measurement of organic molecules fixed on substrate by a simple structure. The current sequence represents a nucleic acid sequence used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Probe for measuring device of intermolecular interaction of organic molecule, comprises probe portion coated with electroconductive film and organic film.
                                                                                                                                                                                                                                                                                                                                                                          Modified nucleic acid used in intermolecular interaction measurement #1.
                                                                                                                                  Gaps
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                                                                                     , DB 1; her.
0. 1e+02;
0; Indels
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1.3%; Score 35; DB 1; Length 40;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 35; Conservative 0; Mismatches 0; Indels
                                                               Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "HS-(CH2)6-0-PO2-0-Thymine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 40 BP; 0 A; 0 C; 0 G; 40 T; 0 U; 0 Other;
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                                                                                                                                                                                                1.3%; Score 35; DB
100.0%; Pred. No. 1e+
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                               Intermolecular interaction; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                BP.
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                                                                                                                                                                                                                                                                                  AED51678 standard; DNA; 40
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Best Local Similarity 100.0
Matches 35; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                                                                                                                                                                  AED51678;
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The new invention relates to a method for detecting (M1) a target gene having a specific sequence in a specimen, by adding a single-stranded comparing a sequence in a specimen, by adding a single-stranded comparing a sequence of the target, therefore forming double-stranded complementary sequence of the target, therefore forming double-stranded complementary sequence of the target, therefore forming double-stranded complementary sequence of the intercalator binds with double-stranded nucleic acid and detecting intercalator bound nucleic acid. Also claimed are a gene detector, which detects a gene having a specific sequence, in a specimen. In (M1), the electrochemical measurement applies voltage with the electrochemical is specific acid is covalently bonded with the compound which as the intercalating agent, is measured. The intercalating agent is specifically inserted in the double-stranded nucleic acid, where the intercalating agent includes a compound which has the double-stranded nucleic acid by the double-stranded nucleic acid by ilght irradiation, the electrochemical active site which has electrochemical activity, and the connection region which connects the councer the interchemical activity, and the connection region and the electrochemical active site. (M1) is useful in gene diagnosis and/or infectious disease diagnosis. The new sequence is a 40mer poly-A detecting DNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting target gene with specific sequence by adding intercalator to double-stranded nucleic acid formed by adding sample having single-stranded target, to probe, performing photo irradiation and detecting intercalator bound nucleic acid.
                                                                                                                                                                                                                                                                                                                     DNA detection; hybridization; diagnosis; infection; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bando K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Murayama R, Tabata J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
                                         40 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; SEQ ID NO 3; 35pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MATU ) MATSUSHITA ELECTRIC IND CO LTD. (MICR-) MICROTEC CO LTD.
                                                                                                                                                                                                                                                                               40mer poly-A detecting DNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hori J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-APR-2005; 2005WO-JP008374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-APR-2004; 2004JP-00128940
                                                                                                                                                 AED68744 standard; DNA; 40
                                                                                                                                                                                                                                   12-JAN-2006 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Maeda M, Akimoto K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2005-769097/78.
                                                                                                                                                                                                                                                                                                                                                  biochip; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                               WO2005103695-A1.
                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-NOV-2005
                                                                                                                                                                                           AED68744;
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Gene detection method for detecting genes having specific sequences, involves adding sample to electrode having probe, hybridizing sample and probe, adding intercalating agent, performing light irradiation and detecting agents forming bonds.

Egashira N;

Bando K,

Tabata J,

Murayama R,

Hori J,

WPI; 2006-090483/09.

(MATU ) MATSUSHITA ELECTRIC IND CO LTD.

06-JUL-2004; 2004JP-00199157. 30-JUN-2005; 2005WO-JP012080.

Specifically

invention relates to a gene detection method.

The new

Example 1; Page 25; 40pp; Japanese.

DNA detection; diagnosis; infection; antimicrobial; drug discovery; ss.

AEF24436 standard; DNA; 40 BP.

(first entry)

09-MAR-2006

AEF24436;

40mer poly A DNA sequence.

WO2006003991-A1. Unidentified

12-JAN-2006.

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The new invention relates to a gene detection method. Specifically described is a method of detecting single stranded probes on an electrode, described is a method of detecting single stranded probes on an electrode, adding an intercalating agent to the electrode, hybridizing the sample with the probe to comparison acarying out light irradiation for covalent bonds forming between nucleic acid and agent and detecting agent forming a covalent comprising compounds represented by "Lb-Ib and "Lc-Fb at each of the sites, where Fa and Fb are electrochemically active site having substituent (s) comprising compounds represented by "Lb-Ib and "Lc-Fb at each of the sites, where Fa and Fb are electrochemically active site having substituent (s) comprising compounds represented by "Lb-Ib and "Lc-Fb at each of the sites, where Fa and Fb are electrochemically active site having substituent (s) binding site which is to be inserted specifically into the double comprises acid und is capable of forming a covalent bond with the couple stranded nucleic acid und is capable of forming a covalent bond with the couple stranded nucleic acid und is capable of forming a covalent bond with the computions sites that links the double stranded nucleic acid und is capable of comprises applying a voltage with respect to the electrochemically a compounds. The detection process comprises applying a voltage with respect to the electrochemical light-emission quantity by the covalent bonded double stranded nucleic acid and intercalating agent. The compound that has photosensitivity is a furocoumarin derivative, preferably a psoralen derivative. (MI) is useful in gene diagnosis, infectious diseases confined in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published pct sequences. The present sequence is a 40-
mer poly A DNA sequence related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. No. 1e+02; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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2709 ААААААААААААААААААААААААААААААА 

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Gaps

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Query Match 1.3%; Score 35; DB 1; Length 40; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 35; Conservative 0; Mismatches 0; Indels

Best Loca Matches

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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743 

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The invention relates to a method for detecting a gene with a specific sequence. The method involves: (a) hybridizing a target nucleic acid to a sequence. The method involves: (a) hybridizing a target nucleic acid to a probe immobilized on an electrode; (b) contacting the resulting duplex with a photoactivatable intercalating agent that has an electrochemical active site; (c) irradiating the complex so that the intercalating agent complex is a covalent bond to the DNA duplex; and (d) detecting the DNA duplex; of an electrochemical measurement (especially electrochemical light emission). The method of the invention is useful for detecting a gene of genese diagnosis and genome based drug discovery. By covalently attaching the intercalating agent to the DNA duplex, the background noise associated with non-specific adsorption of an intercalating agent to double-tranded DNA is reduced, making the method highly sensitive. The present sequence represents a polya (mer used as a control target sequence) and control target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting gene with specific sequence, by denaturing gene sample into single-stranded molecule, reacting single-stranded nucleic acid with nucleic acid probe immobilized on electrode, and electrochemically detecting bound nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Egashira N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40 BP; 40 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
AEF05825) in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bando K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA detection; hybridization; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 3; 15pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MATU ) MATSUSHITA DENKI SANGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                              PolyA target sequence, SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tabata N,
                                                                                                                                                                                                   ВР
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                                                                                                                                                                                                   AEF05826 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2006-113221/12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                         RESULT 41
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IID AEFO

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AC AEFO

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AC AEFO

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XXX

AC AEFO

BE POLJ

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1.3%; Score 35; DB 1; Length 40;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
                                       Conservative
                      Best Local Similarity
Matches 35; Conserv
        Query Match
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Gaps

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Score 35; DB 1; Length 41; Pred. No. 1e+02; 0; Mismatches 0; Indels

1.3%; Scc... 100.0%; Pred. No. ... ... 0; Mismatches

35; Conservative

Matches

Query Match Best Local Similarity

2709 ААААААААААААААААААААААААААААААААААА

Bb.

ADO41099 standard; cDNA; 41

RESULT 43 ADO41099

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## 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743

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ss; probe; human; bacteria; virus; prion; parasite; fungus; drug; allergen; influenza; malaria; yellow fever; multiple sclerosia; Alzheimer's disease; lung cancer; breast cancer; stomach cancer.

WO2004046382-A2.

03-JUN-2004

Homo sapiens

Human cDNA probe useful for disease diagnosis #250.

26-AUG-2004 (first entry)

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ADO41099;

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ADN35262/c
ID ADN35262 Btandard; DNA; 41
                                        (first entry)
                                        01-JUL-2004
                        ADN35262;
        SXXXE
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RESULT 42

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The present invention relates to detecting a probe located and/or a target capable of binding specifically to the probe on a substrate comprises the preparation of a substrate with the probe and/or the target for specific binding to the probe located on its surface, and measurement of the substrate surface by time- of-flight secondary-ion mass spectrometry with labeling. The method is for analyzing probe supports or nucleic acids on nucleic acid chips with detection and quantitation of probe conditions and hybrid of probe with target nucleic acid, which is applicable in gene analysis, disease diagnosis and species identification. The present sequence represents a probe sequence used for hybridization tests.
                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of probe supports or nucleic acids on nucleic acid chips by halogen-based time-of-flight secondary-ion mass spectrometry, applicable in gene analysis, disease diagnosis and species identification.
                                      secondary-ion mass spectrometry; gene analysis; disease diagnosis; species identification; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 41 BP; 0 A; 0 C; 0 G; 38 T; 3 U; 0 Other;
         Probe sequence used for hybridization tests #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; SEQ ID NO 7; 68pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                 Ï
                                                                                                                                                                                                                                                                                                                                                 Hashimoto
                                                                                                                                                                                                                                       28-JUN-2002; 2002JP-00190010.
28-JUN-2002; 2002JP-00191391.
28-JUN-2002; 2002JP-00191414.
                                                                                                                                                                                                          26-JUN-2003; 2003WO-JP008104.
                                                                                                                                                                                                                                                                                                                                                 Takase H,
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-203385/19.
                                                                                                                                                                                                                                                                                                                (CANO ) CANON KK.
                                                                                                                                      WO2004003532-A1.
                                                                                                                                                                        08-JAN-2004
                                                                                                                                                                                                                                                                                                                                                     Okamoto T,
                                                                                                      Synthetic.
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The invention relates to an improved multiplex ligation-dependent amplification method for detecting specific single stranded target cucleic acids in samples using a plurality of probe sets comprising a tucleic acids in samples using a plurality of probe sets comprising a test 2 probes. Each probe comprises a target specific region and a noncomplementary region comprising a primer binding site. The probes in each set are ligated when hybridised to a target nucleic acid and amplified by a primer set. The method is used for detecting a nucleotide polymorphism, especially a single nucleotide polymorphism; detecting multiple single stranded target nucleic acids sequences (through the detection of multiple single stranded nucleic acids in a sample; and detection of a break point region in rearranged nucleic acids. By using a femtomolar amount of the probes, a large number of different probe sets can be used to simultaneously detect and quantify a corresponding large number of target sequences with high specificity. The present DNA sequence is biotin-dial labelled fluorescent oligonucleotide which is used for the hybridisation
                                                                                                                                                                                                                                                                                                                                        An improved multiplex ligation-dependent amplification method for detecting specific single stranded target nucleic acids in samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NS5B; replicon; hepatitis C virus infection; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 43 BP; 0 A; 0 C; 0 G; 43 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.3%; Score 35; DB 1; Lot 100.0%; Pred. No. 1.1e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC.
(RICE-) IST RICERCHE BIOL MOLECOLARE ANGELETTI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NS5B genotype 2b RNA recovery primer, dA(34).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    43 АААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         selection reaction of human mRNA samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hepatotropic; virucide; ss; PCR; primer
                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Page 137; 158pp; English.
                                                                                                                                  15-FEB-2000; 2000EP-00200506.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEA32851 standard; DNA; 34 BP
                                                                                  15-FEB-2001; 2001WO-EP001739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-NOV-2004; 2004WO-US036575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-NOV-2003; 2003US-0517605P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            35; Conservative
                                                                                                                                                                                     (SCHO/) SCHOUTEN J P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                        WPI; 2001-550053/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2005047463-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-2005
                                  23-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-2005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEA32851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AEA32851
          셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising at least 10 oligonucleotides chosen from oligonucleotide from specification. (I) comprising pl is useful for determining gene capters of a cise of probes informative for disease diagnosis, as given in the specification. (I) comprising pl is useful for determining gene capturers characteristic of a disease or condition or its stage in an corganism, for preparing a test gene transcript pattern, for diagnosing or identifying or monitoring a disease or condition or its stage in an corganism. (I) is useful in diagnosing or identifying or monitoring any condition, aliment, disease or reaction that leads to the relative increase or decrease in the activity of information genes of any organism regardless whether the changes caused by the influence of bacteria, virus, prions, parasites, fungi, drugs or allergens, where the disease or cancer such as lung cancer, breast cancer and stomach cancer. (I) enables analysis of gene expression within cells which provides individual from which the cells are derived. (I) enables early detection of a disease or condition or its stage after the onset of the disease or condition, even years before other subjective or objective symptoms appear. (I) enables prevention of the possibility of poor analysis, e.g., represents a human cDNA probe useful for disease diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mRNA hybridisation selection reaction biotin-dT3 oligonucleotide.
                                                                                                                                                                                                                                                                                Set of oligonucleotide probes, useful for diagnosing breast cancer or Alzheimer's disease, comprising specific number of oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, multiplex ligation-dependent amplification, amplicon, single nucleotide polymorphism, hybridisation selection reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                           invention relates to a set (I) of oligonucleotide probes (P1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 41 BP; 41 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note = "Biotin-labelled Thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 АААААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 648; 301pp; English.
                                                                                                                                                                                Loenneborg A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/mod base= OTHER
21-NOV-2003; 2003WO-GB005102.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD17216 standard; DNA; 43 BP
                                                21-NOV-2002; 2002GB-00027238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                Sharma P, Sahni NS,
                                                                                                  (DIAG-) DIAGENIC AS.
                                                                                                                                                                                                                             WPI; 2004-420641/39.
                                                                                                                          (JONE/) JONES E L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200161033-A2
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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AAD17216;

AAD17216/c RESULT 44

Query Match Best Loca Matches

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Gaps

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Length 43; 0; Indels

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The invention relates to a novel method for enhancing the ability of a genotype 2b NSBB sequence to function in a replicon. The method comprises altering either or both the genotype 1b NSBB sequence to encode one or more adaptive mutations, or a genotype 1b NSBB sequence to encode one or adaptive mutations, or a genotype 1b NSBB sequence to encode one or adaptive mutations or a genotype 1b NSBB sequence to encode one or cald (ARB32874) sequence given in the specification. The invention in form with the specification. The invention of a further comprising a given in the specification, The invention of the sequence of a hepatitis C virus (HCV) replicon encoding a fully defined 1394 amino acid (ARB32849) as genotype 2b NSSB encoding nucleic acid (ARB32849) as equence of a genotype 1b replicon, where NSBB contains a Val-218-Ala modification, and substantially all of a genotype contains a Val-218-Ala modification, and substantially all of a genotype C a replicon of one of the method sor chimeric replicon, where the replicon is expressed in the cell. The method has virucide activity. The method is useful for enhancing the ability of a genotype 2b NSSB sequence to the setul for measuring the ability of a genotype 2b NSSB sequence to useful for measuring the ability of a compound to inhibit replicon contains a replicon. The chimeric replicon and recombinant cell are useful for measuring the ability of a compound to inhibit the onset of hepatitis C virus (HCV) infection in a patient. The method is useful convellence represents a primer used in the exemplification of the novel method of the invention.
                                                                                                 Enhancing ability of genotype 2b NS5B sequence to function in replicon for producing replicons containing functional genotype 2b NS5B, and measuring ability of compound to inhibit replicon activity, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                non-structural protein; NS5B; pharmaceutical;
hepatitis C virus infection; antiinflammatory; hepatotropic; virucide;
gastrointestinal disease; infection; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reverse primer used in PCR1 to amplify NS5B RNA from HCV genotypes.
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Σ
  Pizzuti
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1.1e+02;
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  Flores OA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 34 BP; 34 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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nes 34; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.2%; Score 34;
  Lafemina RL,
                                                                                                                                                                                                             Example 1; SEQ ID NO 5; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEB86391 standard; DNA; 34 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-JAN-2004; 2004US-0535708P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JAN-2005; 2005WO-US000292
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    Graham DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MERI ) MERCK & CO INC.
                                                                                                                                                                    treating hepatitis C.
                                                                WPI; 2005-372359/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2005070957-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-2005
      Ludmerer SW,
                            Traboni C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEB86391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 46
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Novel replicase complex comprising hepatitis C virus NSSB replicase, a 3 nucleotide-long template to which a 2 nucleotide-long primer is annealed, and template and primer which do not form a stable duplex in the absence
                                                                                                                        The specification describes non-structural protein NSSB from clinically important hepatitis C virus (HCV) genotypes. NSSB is a RNA-dependent RNA bolymerase. NSSB polypeptides and polymucleotides are useful for evaluating inhibitors of NSSB. These inhibitors may be used to treat HCV infection. PCR primers AEB86390 and AEB86392-AEB86393 were used in a nested PCK reaction to amplify NSSB RNA, with PCR primer AEB86391 as the creverse primer in PCR1. The primers were used for rescue and characterization of NSSB.
                                                       Novel purified hepatitis C virus RNA-dependent RNA polymerase e.g. NS5B polypeptide, useful for evaluating ability of its inhibitor.
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ‡
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus (HCV) NS5B replicase RNA synthesis template
     Lafemina RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus; HCV; NS5B replicase; 88; RNA polymerase.
                                                                                                                                                                                                                                                                                Length 34;
                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                    Sequence 34 BP; 34 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                             1.2%; Score 34; DB 1; Le
100.0%; Pred. No. 1.1e+02;
tive 0; Mismatches 0;
        Flores OA,
                                                                                                                                                                                                                                                                                                                                    2709 АААААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                              34
                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID NO 12; 39pp; English
        Ludmerer SW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleotide-long template to which and template and primer which do of HCV NS5B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-APR-2001; 2001US-00828034
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK99272 standard; RNA; 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                            34; Conservative
         Simcoe AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-582330/62
                                   WPI; 2005-533995/54.
                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhong W, Hong Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HONG/) HONG Z.
(FERR/) FERRARI E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZHON/) ZHONG W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2002064771-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-2002.
           Graham DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK99272;
                                                                                                                                                                                                                                                                                                                 Matches
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The invention relates to a replicase complex comprising a hepatitis C virus (HCV) NSSB replicase protein, a linear nucleic acid template and a complementary nucleic acid primer which is annealed to the 3' terminus of the template, where the template is at least three nucleotides and the primer is two or three nucleotides, and the template and primer do not form a stable duplex in solution in the absence of the HCV NSSB protein.

Example; Page 6; 17pp; English.

to characterise the biological relevance of lead compounds.

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AAD27116 standard; RNA; 36 BP
                                                                                                                                                                                       99US-00309670
                                                                                                                                                                                               99US-00309670
                                                                                                                                                                                                                        Shong W, Hong Z, Lau JYN;
                                                                                                                                  (first entry)
                                                                   Query Match
Best Local Similarity 100.0
Matches 34; Conservative
                                                                                                                                                                                                                                WPI; 2002-096587/13.
                                                                                                                                                                                                           (HONG/) HONG Z. (LAUJ/) LAU J Y N.
                                                                                                                                                                                                       (ZHON/) ZHONG W.
                                                                                                                                  09-APR-2002
                                                                                                                                                                                               11-MAY-1999;
                                                                                                                                                                                       11-MAY-1999;
                                                                                                                                                               Unidentified
                                                                                                                                                                       US6322966-B1
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                                                     templates
                                                                                                                          AAD27116;
                                                                                                          RESULT 48
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The complex is useful for detecting HCV replicase activity and permits establishment of sensitive RNA-dependent RNA polymerase assays to screen and evaluate antivital inhibitors and to improve the specificity and efficacy of the inhibitors. The complex is also useful in the development of a reliable system for determining kinetic and thermodynamic constants of HCV NSSB-catalysed nucleotide incorporation and investigation of specifically, the short RNA template and primer pairs are useful in screening assays which are used for determining kinetic, thermodynamic and mechanistic properties of NSSB replication and ultimately in the development of inhibitors of NSSB. Newly identified inhibitors of replicase activity may be used for development and ultimately in the selection and account of inhibitors of NSSB. Newly identified inhibitors of replicase activity may be used for development anti-HCV pharmaceuticals. Sequences ABK99271-ABK99296 represent HCV NSSB replicase RNA synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 36 BP; 34 A; 0 C; 2 G; 0 T; 0 U; 0 Other;
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RESULT 49 AAL07487 The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3' end capable of forming a copyback structure, a HCV non-structural protein 58 (NSSB), ATP, GTP, TP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is radiolabelled and an assay buffer that supports replication activity of NSSB. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates, RNA template, AA used to direct RNA synthesis by HCV RNA polymerase. Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B; lead compound; RNA polymerase; ss. Example 1; Fig 1A; 10pp; English.

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optimal RNA templates can be used for developing a system to characterise HCV NSSB polymerase mechanistically and kinetically and for designing small RNA molecules to co-crystallise with HCV NSSB polymerase. The assay system of the invention is useful for detecting HCV replicase activity. The nucleic acid synthesised by NSSB is detected by evaluating an autoradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template, AA used to direct RNA synthesis by RNA polymerase proteins of HCV, BVDV and polivirus. This sequence is used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                   Human; reproductive system related antigen; reproductive system disorder;
                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                           Human reproductive system related antigen DNA SEQ ID NO: 10175.
                                                                                                                                                                                       ;
0
                                                                                                                                                             Score 34; DB 1; Length 36;
Pred. No. 1.1e+02;
                                                                                                                                                                                      0; Indels
                                                                                                                                      Sequence 36 BP; 34 A; 0 C; 2 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                    100.0%; Prec. ...
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2000US-0217496P.
2000US-0218290P.
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2000US-0198123P
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2000US-0216647P.
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2000US-0220964P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0189874P
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14-AUG-2000; 2000US-0225759P
                                                                                                                                                               1.2%;
                                                                                                                                                                                                                                                                                                 AAL07487 standard; DNA; 38
                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                     34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                cancer; gene therapy; ds
                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200155320-A2.
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18-APR-2000;
19-MAY-2000;
07-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-MAR-2000;
16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUN-2000;
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PR 22-AUG-2000; 2000US-022679P.
PR 22-AUG-2000; 2000US-0226681P.
PR 22-AUG-2000; 2000US-0226681P.
PR 22-AUG-2000; 2000US-022944P.
PR 01-SEP-2000; 2000US-022943P.
PR 01-SEP-2000; 2000US-022943P.
PR 01-SEP-2000; 2000US-022943P.
PR 01-SEP-2000; 2000US-022943P.
PR 01-SEP-2000; 2000US-022944P.
PR 01-SEP-2000; 2000US-022943P.
PR 01-SEP-2000; 2000US-02239P.
PR 01-SEP-2000; 2000US-023194P.
PR 01-SEP-2000; 2000US-023194P.
PR 01-SEP-2000; 2000US-023194P.
PR 01-SEP-2000; 2000US-023194P.
PR 14-SEP-2000; 2000US-023194P.
PR 14-SEP-2000; 2000US-023194P.
PR 14-SEP-2000; 2000US-023196P.
PR 21-SEP-2000; 2000US-023196
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Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 10175; 1297pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 38 BP; 34 A; 1 C; 2 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rosen CA, Barash SC, Ruben SM;
      2000US-0249207P.
2000US-0249209P.
2000US-0249210P.
2000US-0249211P.
2000US-0249211P.
2000US-0249211P.
2000US-0249213P.
2000US-0249213P.
2000US-0249214P.
2000US-0249216P.
2000US-0249216P.
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2000US-0249245P.
2000US-0249264P.
2000US-0249265P.
2000US-0249297P.
2000US-0249299P.
2000US-0249299P.
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2000US-0251030P.
2000US-0251988P.
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2000US-0251479P.
2000US-0251856P.
2000US-0251868P.
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2000US-0251989P.
2000US-0251990P.
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                                                                                                                                                                                                                                                                                                                                                             2000US-0254097P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-465570/50.
                                                                                                                        17-NOV-2000;
10-DEC-2000;
05-DEC-2000;
06-DEC-2000;
        17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
                                                                                                              17-NOV-2000;
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ss; probe; human; bacteria; virus; prion; parasite; fungus; drug;

Human cDNA probe useful for disease diagnosis #472

(first entry)

26-AUG-2004

AD041321;

BP.

ADO41321 standard; cDNA; 40

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The invention relates to a set (I) of oligonucleotide probes (PI), comprising at least 10 oligonucleotides chosen from oligonucleotide from comprising at least 10 oligonucleotides chosen from oligonucleotide from comprising at least 10 oligonucleotides chosen from oligonucleotide from competition. (I) comprising PI is useful for determining gene expression pattern of a cell, for preparing a standard gene transcript pattern characteristic of a disease or condition or its stage in an or comparism, for preparing a disease or condition or its stage in an originism. (I) is useful in diagnosing or identifying or monitoring any condition, aliment, disease or reaction that leads to the relative condition, parasites, fundin, the activity of information genes of any organism regardless whether the changes caused by the influence of bacteria, virus, parasites, fundi, drugs or allergens, where the diseases confinence such as lung cancer, breast cancer and stomach cancer. (I) cancer such as lung cancer, breast cancer and stomach cancer. (I) cancer such as lung cancer, breast cancer and stomach cancer. (I) cancer such as lung cancer, breast cancer and stomach cancer. (I) cancer such as lung cancer, breast cancer and stomach cancer. (I) cancer such as nalysis of gene expression within cells which provides information on the state of those cells and importantly the state of the individual from which the cells are derived. (I) enables early detection of a disease or condition, even years before other subjective or objective symptoms appear. (I) enables prevention of the possibility of poor analysis, e.g., represents a human cDNA probe useful for disease diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotide probes, useful for diagnosing breast cancer or Alzheimer's disease, comprising specific number of oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
    allergen; influenza; malaria; yellow fever; multiple sclerosis; Alzheimer's disease; lung cancer; breast cancer; stomach cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2%; Score 33.6; DB 1; Length 40; 10.0%; Pred. No. 1.3e+02; ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphism detection related oligonucleotide SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 40 BP; 31 A; 2 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphism; detection; mass spectroscopy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 1378; 301pp; English.
                                                                                                                                                                                                                                                                                                                                    Loenneborg A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABN89412 standard; DNA; 40 BP
                                                                                                                                                                                      21-NOV-2003; 2003WO-GB005102.
                                                                                                                                                                                                                           21-NOV-2002; 2002GB-00027238
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allergen; influenza;
                                                                                                                                                                                                                                                                                                                                    Sharma P, Sahni NS,
                                                                                                                                                                                                                                                                    (DIAG-) DIAGENIC AS. (JONE/) JONES E L.
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-420641/39
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                                                                                                     WO2004046382-A2.
                                                                Homo sapiens
                                                                                                                                             03-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-AUG-2002
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DNA by: (a) preparing a DNA sample from patients containing the DNA region in which the target polymorphism is located; (b) hybridising to an appropriate oligonucleotide fragment, immobilised on a support; and (c) detecting the hybridised target DNA by mass spectroscopy. The method can be used for the investigation and diagnosis of gene-related diseases. The method allows polymorphisms to be detected rapidly and effectively in a large number of specimens. The present sequence represents an oligonucleotide which is used in an example from the present invention
                                                                                                                                                                                                                                    mass spectroscopy for investigation and
                                                                                                                                                                                                                                                                                                          The present invention describes a method for detecting polymorphisms in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kit for the detection and typing of small round-structured virus (SRSV) strains for investigation of food poisoning outbreaks, contains
                                                                                                                                                              Morikawa M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Small round structured virus; SRSV; food poisoning; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2%; Score 33.2; DB 1; Length 40; 92.1%; Pred. No. 1.4e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                           Kimura N, Nagashima R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer SEQ ID 24 used to amplify SRSV specific cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sato T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2701 TITGTACTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 40 BP; 30 A; 0 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kamata K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NINA-) JAPAN NAT INST INFECTIOUS DISEASES (DENK-) DENKA SEIKEN KK.
                                                                                                                                                                                                                                 Detection of DNA polymorphism by me diagnosis of gene-related diseases.
                                                                                                                                                                                                                                                                             Example 3; Page 28; 34pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Miyamura T,
                                                                                                                                                           Nakajima K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF29153 standard; DNA; 33 BP.
                                                                      12-DEC-2001; 2001WO-JP010892.
                                                                                                   12-DEC-2000; 2000JP-00378091,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Small round structured virus
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                                                                                                                             (CHUS ) CHUGAI SEIYAKU KK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 92.1
                                                                                                                                                                                                      WPI; 2002-508814/54.
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                                                                                                                                                           Tamiya G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-080848/09.
             WO200250307-A1.
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                                          27-JUN-2002.
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                                                                                                                                                           Inoko H, ?
Okamoto K;
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WPI; 2004-727850/71.

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                                  This invention relates to a kit for the detection and typing of small round structured virus (SRSV) strains. The kit contains antibodies directed against peptides represented in sequences AAB49700 - AAB49710, which are each SRSV strain specific. Polynucleotide sequences AAF20141 - AAF20151 represent cDNA encoding the strain specific proteins. The kit is sueed for detecting and typing strains of SRSV in order to prevent the spread of infection and to examine the epidemiology of outbreaks. PCR primers AAF29152 - AAF29163 are used to amplify SRSV strain specific CDNA
                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                             Multisignal labeling reagent associated oligonucleotide seqid 1.
                                                                                                                                                                                                                                                                                                                               labeling molecule; solubility; multisignal labeling reagent; ss; DNA-RNA hybrid.
                                                                                                                                                                     ;
0
                                                                                                                                                    DB 1; Length 33;
1.3e+02;
hes 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag= e
note= "Allylamine modified uridine"
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/note= "Allylamine modified uridine"
                                                                                                                                                                                                                                                                                                                                                                                                     a
"Allylamine modified uridine"
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"Allylamine modified uridine"
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"Allylamine modified uridine'
                                                                                                                                                                                                                                                                                                                                                                                                                                       "Allylamine modified uridine"
                                                                                                                                 Sequence 33 BP; 0 A; 0 C; 0 G; 33 T; 0 U; 0 Other;
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                                                                                                                                                                                                   Donegan JJ;
                                                                                                                                                    1.2%; Score 33; DB 100.0%; Pred. No. 1.3 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                 cocation/Qualifiers
                    Example 1; Page 75; 84pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stavrianopoulos JG,
                                                                                                                                                                                                                                                          ADS19106 standard; DNA; 33 BP
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STAVRIANOPOULOS J G.
DONEGAN J J.
                                                                                                                                                                                                                                                                                               30-DEC-2004 (first entry)
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The invention describes a composition (1) of matter comprising an oligomer or polymer having two or more labeled groups, where the label or oligomer or polymer, one or more labels are chemically linked to the oligomer or polymer, one or more reactive groups, and one or more charged groups where the charged groups are covalently linked to the oligomer or polymer or comprise part of the backbone of the oligomer or polymer, or any of their combination. Also described are: a composition (II) comprising a target molecule that has been labeled using (I); and a composition (III) prepared by a target of labeling reagent having the formula (FI) or (F2), (II) reacting the target and the labeling reagent to form the composition having the correcting or guantifying an analyte in a specimen; and detecting or quantifying an analyte in a specimen; and detecting or quantifying an analyte in a specimen; and detecting or contantifying an analyte specific moiety, contacting the contantify the amount of (III) or (III) is useful for (III) or (III) with a specimen suspected of containing the analyte, and measuring the amount of (III) or (III) bound to analytes in the specimen contacting the analyte with sensitivity. In (I), the multiple labeled groups increases the amount of signal that is added to the analyte specific moiety, the amount of signal that is added to the analyte of charged groups increases the groups to a desirable target and the presence of charged group increases contacting or account of containing the multiple labeled groups to a desirable target and the presence of charged group increases
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Composition of multi signal labeling reagents, useful for detecting or quantifying analyte in specimen, has oligomer/polymer having labeled moieties, reactive groups and charged groups linked to oligomer/polymer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus; HCV; NS5B replicase; 88; RNA polymerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.2%; Score 33; DB 1; L. 100.0%; Pred. No. 1.3e+02; tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 1; 20pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK99273 standard; RNA; 36
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HONG Z.
FERRARI E.
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WPI; 2002-582330/62. 11-MAY-1999; Query Match Matches RESULT 55 ઠે d 

The invention relates to a replicase complex comprising a hepatitis C virus (HCV) NSSB replicase protein, a linear nucleic acid template and a complementary nucleic acid primer which is annealed to the 3' terminus of the template, where the template is at least three nucleotides and the complementary nucleic acid primer which is annealed to the 3' terminus of the template, where the template is at least three nucleotides and the complex is two or three nucleotides, and the template and primer do not complex is useful for detecting HCV replicase activity and permits catablishment of sensitive RNA-dependent RNA polymerase assays to screen and evaluate antiviral inhibitors and to improve the specificity and efficacy of the inhibitors. The complex is also useful in the development of a reliable system for determining kinetic and thermodynamic constants of HCV MSSB-catalysed nucleotide incorporation and investigation of mechanistic inhibitors for mis-incorporation and ultimately in the specifically, the short RNA template and primer pairs are useful in screening assays which are used for determining kinetic, thermodynamic development of inhibitors of NSSB replication and ultimately in the catablicase activity may be used for developing anti-HCV pharmaceuticals. Novel replicase complex comprising hepatitis C virus NSSB replicase, a 3 nucleotide-long template to which a 2 nucleotide-long primer is annealed, and template and primer which do not form a stable duplex in the absence of HCV NSSB. Example; Page 6; 17pp; English templates

Sequence 36 BP; 33 A; 0 C; 2 G; 0 T; 1 U; 0 Other;

; 0 1.2%; Score 33; DB 1; Length 36; 100.0%; Pred. No. 1.3e+02; tive 0; Mismatches 0; Indels 33; Conservative Best Local Similarity

2709 ААААААААААААААААААААААААААААА 

AAD27117 standard; RNA; 36 09-APR-2002 (first entry) AAD27117;

RNA template, AU used to direct RNA synthesis by HCV RNA polymerase.

Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B; lead compound; RNA polymerase; ss.

Unidentified

US6322966-B1

27-NOV-2001.

99US-00309670

99US-00309670 11-MAY-1999;

(ZHON/) ZHONG W. (HONG/) HONG Z. (LAUJ/) LAU J Y N.

Shong W, Hong Z, Lau JYN;

WPI; 2002-096587/13.

template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates Assay system for hepatitis C virus replicase activity comprises RNA

Example 1; Fig 1A; 10pp; English.

The present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3' end capable of forming a copyback structure, a HCV non-structural protein 5B (NSSB), ATP, GTP, CTP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is caid-labelled and an assay buffer that supports replication activity of NSSB. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds or to characterise the biological relevance of lead compounds. The coptimal RNA templates can be used for developing a system to characterise HCV NSSB polymerase mechanistically and kinetically and for designing CT from the invention is useful for developing a system to characterise correction products separated by gel electrophoresis. The structuradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template, AU used to direct RNA synthesis by RNA copymerase proteins of HCV, BVDV and polivirus. This sequence is used in the exemplification of the invention 

Sequence 36 BP; 33 A; 0 C; 2 G; 0 T; 1 U; 0 Other;

Gaps ö 1.2%; Score 33; DB 1; Length 36; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels 33; Conservative Similarity Query Match Local Matches

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2709 ААААААААААААААААААААААААААААА 3 АААААААААААААААААААААААААААААА 용

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Gaps

AAD27125 standard; RNA; 37 RESULT 56 AAD27125

AAD27125;

BP.

(first entry) 09-APR-2002 RNA template, (AU)2 used to direct RNA synthesis by HCV RNA polymerase.

Hepatitis C virus; HCV replicase; non-structural protein 5B; NS5B; lead compound; RNA polymerase; ss.

Unidentified

US6322966-B1

27-NOV-2001.

99US-00309670. 11-MAY-1999;

11-MAY-1999; 99US-00309670. HONG Z. LAU J Y N. (ZHON/) ZHONG W. (HONG/) 

Zhong W, Hong Z, Lau JYN;

WPI; 2002-096587/13.

Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates buffer.

Example 1; Fig 2A; 10pp; English.

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Hey replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3' end capable of forming a copyhac tructure, a HVV non-structural protein SB (NSSB), ATP, GTP, CTP, and UTP nucleoside triphosphates (NTPS), where one of the NTP is radiolabelled and an assay buffer that supports replication activity of radiolabelled and an assay buffer that supports replication activity of SSB. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HVV. This activity can be used to screen for anti-HVV replicase compounds or to characterise the biological relevance of lead compounds. The cort contacterise the biological relevance of lead compounds. The cort to characterise mechanistically and kinetically and for designing system to characterise captement occrystallise with HCV NSSB polymerase. The assay stem of the invention is useful for detecting HCV replicase activity. The nucleic acid synthesised by NSSB is detected by evaluating an autoradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template, (AU) 2 used to direct RNA synthesis by the average of the invention of the invention of the invention of the invention and the invention products separated by gel electrophoresis. The average proteins of HCV, BVDV and polivitus. This sequence is used in the average.
                                                                                                                                        the present invention relates to an assay system for hepatitis C virus (HCV) replicase activity. The ages of the control of th
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Gaps
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                             1.2%; Score 32.4; DB 1; Length 37; 97.1%; Pred. No. 1.5e+02; tive 0; Mismatches 1; Indels
Sequence 37 BP; 33 A; 0 C; 2 G; 0 T; 2 U; 0 Other;
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                                                                                                                   33; Conservative
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Sequence of scissile link probe MRC068 (HL). AAN70278 standard; DNA; 32 (first entry) Hybridisation; probe; ss (revised) 03-OCT-2002 26-MAY-1991 EP227976-A. Synthetic. AAN70278; 

86EP-00116906. 85US-00805279 Crosby (MEIO-) MEIOGENICS INC. WPI; 1987-186567/27. Bender R, 04-DEC-1986; 05-DEC-1985; 08-JUL-1987 <u>п</u> Duck

Robertson JG; 3 Synthetic nucleic acid probes - comprising two nucleic acid sequences linked by a scissile linkage.

The patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S-- a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or NNA sequences in a test soln. The acissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Example, p29; 46pp; English.

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Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing
                                                                                                                                                                                                      Gaps
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100.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 0; Indels
                                                                                                                     Sequence 32 BP; 0 A; 0 C; 0 G; 24 T; 8 U; 0 Other;
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Matches 32, Conservative
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a "deoxyribonucleotides." /note= "deoxyribonucleotides." Probe MRCO68; solid support; ribonuclease. /\*tag= b /note= "ribonucleotides." Location/Qualifiers AAN92244 standard; DNA; 32 (first entry) 43. .32 /\*tag= c /\*tag= note= (revised) (revised SS probe MRCO68. misc\_feature misc\_feature 25-MAR-2003 misc\_feature 25-APR-1990 Synthetic AAN92244; AAN92244, 셤

Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence. WPI; 1989-339977/46.

88US-00187814 88US-00187814

29-APR-1988; 29-APR-1988;

WO8910415-A 02-NOV-1989 (MEIO-) MEIOGENICS INC

Bender R;

Duck P,

Disclosure; Page 24; 34pp; English.

Probe MRCO68 is bound by a hydrolysable linkage to a solid support at its 3 and. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. The probe system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg RNase H or ExoIII. (Updated on 31-OCT-2002 to add missing OS field.)

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                                                                AAZ98722;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polymerase and dNTPs meessary for repetitively replicating the tandem repeat. The method involves detecting a binding event between first and second binding members, employing a label to determine the occurrence of the binding members, employing a label to determine the occurrence of the binding event. The tandem repeating units are polyT. The method of the invention is useful in detecting an analyte using repetitive extension along a tandem repeat. The extended nucleic acid may be used for detecting a moiety, particularly involved in a binding event employing a reagent. The current sequence represents a template member oligonucleotide containing a polyT tandem repeat that binds to the extendable oligonucleotide given in ADG33444.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting a label comprising employing (as the label) a reagent having a 's extendable terminus hybridized to a tandem repeat template in combination with a DNA polymerase and dNTPs necessary for repetitively replicating the tandem repeat.
                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method for detecting a label, comprising employing (as the label) a reagent having a 3' extendable terminus hybridised to a tandem repeat template in combination with a DNA
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                                                   Length 32;
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                                                                                                                                                                                                                                                                                                  Binding; tandem repeat; label; analyte detection; ss.
                         Sequence 32 BP; 0 A; 0 C; 0 G; 24 T; 8 U; 0 Other;
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100.0%; Pred. No. 1.4e+02;
iive 0; Mismatches 0;
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 to correct PR field.)
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                                                                                                                                                                                                                                                                        Template oligonucleotide #SEQ ID 2.
                                                                                                                                                                                           ADC33445 standard; DNA; 32 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                               21-FEB-2002; 2002US-0359223P.
08-MAY-2002; 2002US-0379360P.
                                                                                                                                                                                                                                             18-DEC-2003 (first entry)
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on 25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-712717/67.
                                                               Local Similarity
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Matches
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ADC33445/c
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This sequence represents a PCR primer used to determine the full length cDNA sequence of the swine vesicular disease virus (SVDV) gene sequence of Taiwan Yu-Li strain (see AAZSBTT). SVDV is the causarive agent of swine vesicular disease, which is very similar to foot and mouth disease. The invention relates to the wild-type Taiwan Yu-Li strain cDNA sequence, and the gene sequences of the mutant SVDV strains N3, H21 and SP7. The mutant SVDV nucleotide sequence can be used in a vaccine for the prophylaxis of swine vesicular disease. The invention also includes a method for differentiating the mutant SVDV nucleotide sequences from the wild type strain of SVDV, coxsackievirus and foot-and-mouth disease virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mutant strains of swine vesicular disease virus (SVDV) used in vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                       PCR primer used for swine vesicular disease virus gene synthesis.
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                                                                                                                                                                                                                       Swine vesicular disease virus; SVDV; swine vesicular disease;
Taiwan Yu-Li strain; foot and mouth disease; coxsackie virus;
differentiation; vaccine; prevent; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40 BP; 1 A; 3 C; 3 G; 33 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2%; Score 32; DB 1; L. 100.0%; Pred. No. 1.7e+02; iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                through the use of polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   40 AAAAAAAAAAAAAAAAAAAAAAAAA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to prevent swine vesicular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Χ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 6; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yang Y, Jeng
AAZ98722 standard; cDNA; 40 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BIOT-) DEV CENT BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                              Swine vesicular disease virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV12483 standard; DNA; 39 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98EP-00306486.
                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-258616/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SXXXXXXXXXXX
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2709 ААДАДАДАДАДАДАДАДАДАДАДАДАДАДА

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WPI; 1999-181062/15.
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ID ADI3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            원
                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents an oligonuclectide used in an example of the present invention. The present invention describes a method for synthesising a selected oligonuclectide (I) having well defined ends. The method comprises: (a) annealing a primer to a single-stranded (ss) circular template to yield a primed circular template, where the template comprises: (i) at least one nucleotide effective to produce a cleavage site in the cligonuclectide multimer; (b) combining the primed circular template with at least two types of nucleotide ffective to produce a cleavage site in the cligonuclectide two types of nucleotide triphosphates and a polymerase enzyme without the addition of auxiliary proteins to yield a ss oligonucleotide multimer complementary to the circular oligonucleotide template, comprising multiple copies of (I); and (c) cleaving the oligonucleotide cultimer at the cleavage site to produce (I) having well defined ends. The method is used for the large-scale synthesis of box and RNA oligomers for use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                      Rolling circle synthesis of oligo:nucleotide(s) - using primed circular template to produce oligonucleotide multimer for cleavage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         therapeutic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Multimer; probe; diagnosis; synthesis; detection; polymerase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.2%; Score 31.8; DB 1; Length 39; 94.3%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use, e.g. as probes and diagnostic agents and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 39 BP; 0 A; 0 C; 3 G; 36 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ааааааааааасаааааааааааасааааааа 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Col 45; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX30019 standard; DNA; 39 BP
                                                                                                                                                             93US-00047860.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-00910632
                                                                                                                    95US-00393439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 94.3<sup>3</sup>
"harres 33; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                      (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Multimer SEQ ID NO:6.
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                                                                                                                                                           15-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-AUG-1998;
                                                                                                                    23-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-FEB-1999
                           US5714320-A
                                                                        03-FEB-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX30019;
                                                                                                                                                                                                                                                 Kool ET;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kool ET;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 62
         ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  X B X B X B X B X B X B X B X B X B X X B
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                                                                                                                                                              The present invention describes a detectably labelled oligonucleotide multimer, comprising multiple contiguous copies of a repeated oligonucleotides. The detectably labelled oligonucleotide multimer is useful for detecting a trarget molecule. Oligonucleotide multimers may be produced in sufficient quantity to be useful for diagnostic and medical applications. The multimers are useful for affinity labelling of proteins, and for signal amplification in highly sensitive affinity acapture and sequence identification applications. The method provides a faster, cheaper and simpler way for large-scale production of DNA and Allogomers and multimers. The incorporation of labels enables the oligonucleotide multimers to be useful in diagnostics and medicine. The present sequence represents an oligonucleotide used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HPV infection; replication; cytostatic; virucide; cervical dysplasia; carcinoma; oral mucosal cancer; laryngeal; vaccine; gene therapy; double internal trans-acting ribozyme; single; dITRz; ITRz; CHOP portion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating human papilloma virus (HPV) infection, e.g. cervical dysplasias or HPV-associated cervical carcinoma, comprises administering to a patient a nucleic acid molecule that inhibits expression associated with HPV replication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3' flanking RNA of IT ribozyme (Rz) from CHOP portion of SNIPAA cassette.
   multiple
detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
New detectably labelled oligonucleotide multimer, comprising contiguous copies of a repeated oligonucleotide - useful for target molecules in diagnosis and medicinal applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 39 BP; 0 A; 0 C; 3 G; 36 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ä
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thiboutot
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 56; 65pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Christensen N,
                                                                                                                              Example 2; Page 41; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0391795P.
2002US-0417997P.
2003US-0449066P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JUN-2003; 2003WO-US020340.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI32816 standard; RNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PENN-) PENN STATE RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNIPAA cassette; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clawson GA, Pan W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-082869/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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14-OCT-2002; 2
21-FEB-2003; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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to Feature Table Key

record issued on 04-NOV-2004 : Correction

Revised

ន្តដូន

Sequence 31 BP; 11 A; 5 C; 7 G; 8 T; 0 U; 0 Other;

41

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Gaps

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ch 1.1%; Score 31; DB 1; Length 31; 1. Similarity 100.0%; Pred. No. 1.6e+02; 31; Conservative 0; Mismatches 0; Indels

Query Match

Local

Matches

8 셤

870 TCAGTAAGATCATAAGCAGTGATCGAGATCT 900 TCAGTAAGATCATAAGCAGTGATCGAGATCT 31

AEE86835 standard; DNA; 31

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The invention relates to a novel method for treating Human papillomavirus (HPV) infection comprising administering to a patient a nucleic acid molecule that inhibite sexpression associated with HPV replication. The method of the invention has cytostatic and virucide applications and may including ervical dysplasia, cervical excrinema, oral mucosal cancer and laryngeal cancer. Furthermore, the method may be utilised during vaccine production and gene_therapy. The current sequence is that of the HPV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecules from the human genome which include polymorphic sites, useful in methods for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a
                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; resequence; genotype; disease; forensic; paternity testing; single nucleotide polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
/standard_name= "single nucleotide polymorphism"
                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                              1.1%; Score 31.2; DB 1; Length 39;
18.9%; Pred. No. 1.8e+02;
.ve 1; Mismatches 3; Indels
                                                                                                                                                                           production and gene therapy. The current sequence is targeted ribozyme (Rz)-related RNA of the invention
                                                                                                                                                                                                                                              Sequence 39 BP; 32 A; 2 C; 2 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                   2704 GTACTAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human single nucleotide polymorphism (SNP) G22P1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lander ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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22-MAY-2000; 2000US-0206129P.
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                                                                                                                                                                                                                                                                                                                          88.9%;
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Matches 32; Conserv
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18-OCT-2001
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                                                                                                                                                                                                                                                                                              Query Match
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DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.

Location/Qualifiers

/note= "OTHER= 3'-terminal Si(O-Me)3"

J= a base= OTHER

/\*tag=

Key modified\_base

Synthetic

/mod

Novel solid phase-related oligonuclectide Oligo-dr31-Si(O-Me)3 #10.

(first entry)

23-FEB-2006

AEE86835;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding
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les 31; Conservative
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GENOVOXX GMBH.
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The invention relates to the identification of nucleic acid molecules (AAI29513-AAI3114) from the human genome which include polymorphic sites which can predispose individuals to disease. Various genes from a number of individuals were resequenced and single nucleotide polymorphisms (SNPs) in these genes discovered. The method is useful for predicting the presence, absence or severity of a particular phenotype or disorder (e.g. diabetes) associated with a particular genotype. The nucleic acids containing the polymorphic sites may be useful in forensics and paternity testing

Claim 1; Page 102; 145pp; English.

particular genotype:

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23-FEB-2006

AEE86830;

Synthetic.

15-DEC-2005

(HENN/) (GENO-)

CHER/)

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This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged and respected contact of the solid phase with high concentrations of labeled reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.
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                                 DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
Novel solid phase-related oligonucleotide Cy3-Oligo-dT31-NH2
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100.0%; Pred. No. 1.6e+02;
ive 0; Mismatches 0; Indels
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/note= "OTHER= 3'-terminal amide"
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(GENO-) GENOVOXX GMBH.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.
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                                                                                                                                                                                                     Novel solid phase-related oligonucleotide Oligo-dT31-NH2 #5.
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Pred. No.
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/mod_base= OTHER
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100.0%;
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ID AEE86829 standard; DNA; 31
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Best Local Similarity 100.
Matches 31; Conservative
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GENOVOXX GMBH.
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RESULT

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This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid
                                                                                                                   Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components.
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1.1%; Score 31; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.6e+02;
Matches 31; Conservative 0; Mismatches 0;
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AEE86845 standard; DNA;
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  (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
(CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                phase of the invention.
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                                                                                                                                                                                                                                                                                                     of solid phase, useful for parallel, optical analysis of many acids, has reduced non-specific binding of labeled components.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel solid phase-related oligonucleotide Oligo-dT31-Si(O-Me)3 #9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
                                           /note= "OTHER= 3'-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 31; DB 1; L. Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 ААААААААААААААААААААААААААА
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%; Sc.
100.0%; Prec
                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 62; 88pp; German.
                                /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/mod_base= OTHER
                                                                                                                             26-MAY-2004; 2004DE-10025745
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Best Local Similarity 100.
Matches 31; Conservative
              *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   phase of the invention.
                                                                                                                                                                                    HENNIG C.
GENOVOXX GMBH.
                                                                                                                                                                                                                                            Cherkasov D, Hennig
                                                                                                                                                                                                                 CHERKASOV D.
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                                                                       DE102004025745-A1.
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modified_base
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                                                                                                                                                                                                                                                                                                      Surface of
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Length 31; 0; Indels 88.

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                                 This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid phase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                              /*tag= b
/mod_base= OTHER
/note= "optionally labeled with an amide group or Si(O-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel solid support surface for parallel analysis of many individual nucleic acids by optical methods. The invention also describes; a) a solid phase in which the surface shows
nucleic acids, has reduced non-specific binding of labeled components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Surface of a solid support, useful for multiple parallel analysis of nucleic acids by optical methods, having low non-specific binding of labeled components.
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                                                                                                                            DB 1; Length 31;
1.6e+02;
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/mod_base= OTHER
/note= "optionally labeled with Cy3"
                                                                                                          Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
                                                                                                                                                                  100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                       DNA detection; DNA sequencing; primer; ss.
                                                                                                                                                                           1.1%; Score 31;
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                                                                                                                                                                                                                                                                                    Oligonucleotide Cy3-Oligo-dT31-NH2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 62; 88pp; German.
                   Disclosure; Page 62; 88pp; German.
                                                                                                                                                                                                                                AEF12155 standard; DNA; 31 BP
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                                                                                                                                               Matches 31; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHERKASOV D.
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                                                                                                                                       Local Similarity
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modified base
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                                                                                                                              Query Match
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cc reduced non-specific binding of labeled components; b) methods for preparing the novel solid support and c) methods of parallel analysis of many nucleic acid by optical methods, using the solid support. The solid support is made of silica, glass, silicon dioxide or silven, is flat and has nucleic acid chains fixed to it, optionally controved the solid phase is preferably part of a device that allows fluid exchange and it is permeable to light in the wavelength callows fluid exchange and it is permeable to light in the wavelength regions 200-400; 202-2000 or 400-800 nm. An external layer of solid support is removed, then the nucleic acid is coupled to it, optionally after attachment of a linker layer. Alternatively, after removing the coupling, optionally after attachment of a linker, and in either case, additional substances (specifically phosphate, sulfate or carboxy-coupling, optionally after attachment of a linker, and in either case, additional substances (specifically phosphate, sulfate or carboxy-coupling, optionally after attachment of synthesis of nucleic acids. Only part of the surface is attachment or synthesis of nucleic acids. Only part of the surface is cattachment or synthesis of nucleic acids. Only part of the surface is or attachment or synthesis of nucleic acids. Only part of the surface is or additional subsequent setpes are done in a liquid phase. The nucleic acids analyzed represent a single population or many different populations and contains 5-50, 20-200 or 50-500 nucleotides. The linker is 1-50 nm long cadeoxytho-nucleoside triphosphates, in which the label is cleavable. Contains 5-50, 20-200 or 15-500 nucleotides. The linker is 1-50 nm long cadeoxytho-nucleoside triphosphates, in which the label is cleavable. Catternatively the nucleic acid sequences that surface is acid sequences that surfa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             so assay sensitivity is improved. This sequence represents an oligonuclectide used to illustrate the method of the invention.
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ive 0; Mismatches
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/mod_base= 5'-Cy3
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/mod_base= 3'-NH2
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dt31-NH2 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                    Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; dna detection; DNA sequencing; DNA amplification; oligo dT31-Si(O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Optical DNA analysis process-related oligonucleotide dT31-Si(O-Me)3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                   Example 5; Page 66; 94pp; German.
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                                                                                                               Cherkasov D, Hennig C;
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                         (CHENN/) CHERKASOV D.
(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
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GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHER/) CHERKASOV D.
                                                                                                                                                           WPI; 2006-185819/20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide Cy3-dt31- which was used in the development of the novel process of the invention.
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                                                                                                                                                                                                                                                                                                                                                      Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88; dna detection; DNA sequencing; DNA amplification; oligo dt31-NH2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Optical DNA analysis process-related oligonucleotide dt31-NH2.
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100.0%; Pred. No..1.6e+02;
rative 0; Mismatches 0; Indels
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31
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/mod_base= 3'-NH2
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                                                                                    26-MAY-2004; 2004DE-10025695.
                                                                                                                                 26-MAY-2004; 2004DE-10025695.
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                                                                                                                                                                                                                                                                 Cherkasov D, Hennig C;
                                                                                                                                                                                          (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                           (CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                        WPI; 2006-185819/20.
DE102004025695-A1
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modified_base
                                           23-FEB-2006.
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Query Match

Matches

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RXFXBXYFTFFFXSSXXXXBXBXBX

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Gaps

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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide d731-51(0 which was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                fluorescent parallel process to analyse nucleic acid chains in sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; dna detection; DNA sequencing; DNA amplification; oligo Cy3-dt31-.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Optical DNA analysis process-related oligonucleotide Cy3-dt31-NH2
                                                                                                                                                                                                                                                                                                                                                                               1.1%; Score 31; DB 1; Length 31; 100.0%; Pred. No. 1.6e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                   the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
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                                                                                    Example 5; Page 66; 94pp; German.
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Best Local Similarity 100.0
Matches 31, Conservative
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(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
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WPI; 2006-185819/20.
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                                  Optical
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present development of the novel process of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; dna detection; DNA sequencing; DNA amplification; oligo dt31-NH2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Optical DNA analysis process-related oligonucleotide dt31-NH2.
                                                                                                                                                                                                                                                    / Match 1.1%; Score 31; DB 1; Length 31; Local Similarity 100.0%; Pred. No. 1.6e+02; Ale 31; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                              Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
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/mod_base= 3'-NH2
            Example 5; Page 67; 95pp; German.
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(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their

Example 5; Page 67; 95pp; German.

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Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;

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co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present sequence is that of oligonucleotide dt31-NH2 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present sequence is that of oligonucleotide d731-Si (0 which was used in the development of the novel process of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; dna detection; DNA sequencing; DNA amplification; oligo dT31-Si(O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Optical DNA analysis process-related oligonucleotide dT31-Si(O-Me)3.
                                                                                                                                                                                         Length 31;
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                                                                                                                                                  Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
                                                                                                                                                                                  1.1%; Score 31; ....,
100.0%; Pred. No. 1.6e+02;
Hive 0; Mismatches 0;
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Best Local Similarity 100.
Matches 31; Conservative
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(GENO-) GENOVOXX GMBH.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; dna detection; DNA sequencing; DNA amplification; oligo dt31-NH2.
                                                                                 Gaps
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                  Length 31;
                                                                              0; Indels
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Pred. No. 1.6e+02;
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                  Score 31; DB 1; L
Pred. No. 1.6e+02;
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1.1%; Sco. 100.0%; Pred. No. 100.0%; Pred. No. 100.0%; Pred. No. 100.00; Pred. No. 1
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                                                                              Conservative
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GENOVOXX GMBH.
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               Query Match
Best Local Similarity
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-APR-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-FEB-2006.
                                                                              31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                            AEF94718;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                              Matches
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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2739

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AEF94717;

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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dT31-Si(O which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                                                   88; dna detection; DNA sequencing; DNA amplification; oligo dT31-Si(O.
                                                                                                Optical DNA analysis process-related oligonucleotide dT31-Si(O-Me)3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2709 АДАДАДАДАДАДАДАДАДАДАДАДАДАДАД
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                         /mod_base= 3'-Si(0-Me)3
                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 95; 141pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEF94717 standard; DNA; 31 BP
                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAY-2004; 2004DE-10025696
                                                                                                                                                                                                                                                                                                                                                                               26-MAY-2004; 2004DE-10025696
AEF94723 standard; DNA; 31
                                                                 (first entry)
                                                                                                                                                                                                                                                         *tag= b
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Matches 31, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cherkasov D, Hennig C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               HENNIG C.
GENOVOXX GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2006-185820/20
                                                                                                                                                                                                                                                                                                           DE102004025696-A1.
                                                                                                                                                                                                                         Key
modified_base
                                                                                                                                                                     Unidentified
                                                                20-APR-2006
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                                                                                                                                                                                         Synthetic.
                                 AEF94723;
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(GENO-)
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide Cy3-dt31-which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                 ss; dna detection; DNA sequencing; DNA amplification; oligo Cy3-dt31-.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                   Optical DNA analysis process-related oligonucleotide Cy3-dt31-NH2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 31 BP; 0 A; 0 C; 0 G; 31 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 95; 141pp; German.
                                                                                                                                                                                                        /*tag= b
/mod_base= 5'-Cy3
                                                                                                                                                                                                                                                                    /mod_base= 3'-NH2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Baeuml
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Best Local Similarity 100...
These 31; Conservative
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                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hennig C,
                                                                                                                                                                                                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                      (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                        CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2006-185820/20.
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                                                                                                                                                                               Key
modified_base
                                                                                                                                                                                                                                         modified base
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                                                                                                                                Unidentified
                                          20-APR-2006
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                                                                                                                                                   Synthetic.
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Gaps

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Length 31; Indels

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                                                                                                                                                                                                                                                                                                                                       The specification describes a method for determining the presence of severe acute respiratory syndrome coronavirus (SARS-CoV) in a test sample. The method comprises contacting a test sample with a probe and determining whether the hybrid is present in the test sample as an indication of the presence of SARS-CoV in the test sample. The method is useful for distinguishing the presence of SARS-CoV from that of human coronavirus strains 229 and OC43. The method is useful for diagnosing SARS or for monitoring the therapeutic treatment of a SARS-CoV-infected individual. The detection probes of the invention are useful as amplification oligonucleotides or helper oligonucleotides. The present sequence represents a homopolymer tail with a flexible linker for use
                   severe acute respiratory syndrome; SARS; SARS-CoV; human coronavirus; human coronavirus strain 229B; human coronavirus strain OC43; SARS infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
        Homopolymer tail with flexible linker for use with a capture probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                            Vijaysri S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%; Score 31; DB 1; Length 33; 00.0%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trichomonas vaginalis nucleic acid target capture probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 33 BP; 30 A; 0 C; 0 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                           Getman DK,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TAAAAAAAAAAAAAAAAAAAAAAAAAAA 33
                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 39; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                with a capture probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; detection; diagnosis.
                                                                                                                                                                                                                           Nelson NC,
                                                                                                                                                                      25-APR-2003; 2003US-0465428P.
09-MAY-2003; 2003US-0469294P.
                                                                                                                                       16-APR-2004; 2004WO-US011636
                                                                                                                                                            2003US-0464049P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADU83547 standard; DNA; 33
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Matches 31; Conservative
                                                                                                                                                                                                                          Kacian DL,
                                                                                                                                                                                                     (GENP-) GEN-PROBE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trichomonas vaginalis
                                                                                                                                                                                                                                              WPI; 2004-795575/78.
                                                                                            WO2004094675-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JS2004235138-A1
                                                                                                                                                            17-APR-2003;
                                                                                                                  04-NOV-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-NOV-2004
                                                                                                                                                                                                                          Linnen JM,
                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                   formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADU83547;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 82
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New detection probe, useful for determining or screening for the presence of Trichomonas vaginalis in a biological sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a detection probe, for determining the presence of Trichomonas vaginalis in a test sample, the probe being up to 100 bases in length and comprising a target binding region which forms a hybrid stable for detection with a sequence contained within any of the [6 (first) target sequences of 26-32 bp under stringent hybridization conditions, where the probe does not form a hybrid stable for detection with nucleic acid derived from Trichomonas tenax under the cited conditions. The detection probe, oligonucleotide, composition, methods, and kits are useful for determining the presence of T. vaginalis in a test sample. This sequence corresponds to a target capture probe for the T. vaginalis DNA and used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA detection, diagnosis, prognosis, DNA amplification, prostatic cancer, cytostatic, PCA3; probe, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31; DB 1; Length 33;
Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 33 BP; 30 A; 0 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prostate cancer specific PCA3 probe SEQ ID NO 41.
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100.0%; Pred. No. 1...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYME-) UNIV MEDICAL CENT NIJMEGEN
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                                                      19-MAY-2003; 2003US-0472028P.
18-MAY-2004; 2004US-00848922.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31; Conservative
                                                                                                                                                                                              Bungo JJ;
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                                                                                                                                                                                                                                               WPI; 2004-821327/81.
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                                                                                                          WEISBURG W
BUNGO J J.
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                                                                                                                                                                                              Weisburg WG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schalken JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-APR-2005
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                                                                                                          (WEIS/)
(BUNG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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The invention describes diagnosing or prognosing prostate cancer in a patient comprising amplifying a prostate cancer specific PCA3 RNA using a patient comprising and detecting an amplification product derived from it, where the amplification product is associated with a presence of prostate cancer or predisposition to prostate cancer in the patient. Also described is a diagnostic kit comprising a first container containing a first pair of primers designed to amplify a PCA3 RNA across an exon junction of a PCA3 gene. The method and kit are useful for diagnosing or prognosing prostate cancer in a patient. This sequence represents a probe used to detect an exon-exon junction to detect PCA3 mRNA

Sequence 33 BP; 30 A; 0 C; 0 G; 3 T; 0 U; 0 Other;

Gaps ; DB 1; Length 33; 1.7e+02; 0; Indels 1.1%; Scor. V 100.0%; Pred. No. 1... O; Mismatches Query Match Best Local Similarity 100.' Marches 31, Conservative

2708 ТАРАРАРАРАРАРАРАРАРАРАРАРАРАРАРА з таарарарарарарарарарарарарара

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AAT93827 standard; DNA; 34 BP AAT93827; RESULT 84 AAT93827 

Antitumoural phosphodiester oligonucleotide 17 with cytotoxic activity. (first entry) (revised) 25-MAR-2003 24-FEB-1998

Phosphodiester; selective binding; cell viability; growth; tumoural cell line; cytotoxic activity; tumour cell; lymphoma; lymphoblastic tumour; ss.

Synthetic.

/note= "phosphodiester oligonucleotide" Location/Qualifiers 1. .34 /\*tag= Key modified\_base

WO9720924-A1

12-JUN-1997

96WO-EP005388 04-DEC-1996; 95IT-MI002539 04-DEC-1995;

(SAIC-) SAICOM SRL.

Quadrifoglio F; Scaggiante B,

WPI; 1997-319771/29.

New phospho:di:esteric oligo:nucleotide(s) - which exert a specific and selective cytotoxic effect on tumour cells, for treating both solid and Claim 10; Page 6; 38pp; English. liquid tumours.

Novel phosphodiesteric oligonucleotides AAT93811-27 are based on the generic formula, in the 3'-5' or 5'-3' direction: (GaTa')a''-(GDTb')b''- (GGTc')a''-(GTTd')d''-(GGTe')a''-(GTTd')d''-(GTTG')a''-where: N and N' = T or G, equal or different from each other; x = 0-8, equal or different from each other; a, b, c, d, e, f, and g = 0-10, equal or different from each other; a', b', c', d', e', f', and g' = 0-30, equal or different from each other; a', b', c', d', e', f', and g' = 1-16, equal or different from each other; a', b', c', d', e', f', and g' = 1-16, equal or different from each other; The oligonucleotides are believed

to selectively bind and sequester some proteins which are essential to the viability and growth of tumoural cell line. They have specific and selective cytotoxic activity against tumour cells, and can be used for treating tumours of the liquid type, in particular of lymphoblastic origin, and of solid type, in particular lymphomas. The present phosphodiester oligonuclectide, at a concentration of 15 micromolar, reduced growth of CCRF-CEM tumoural cells by 71%, which is detectable 48 hours after administration. (Updated on 25-MAR-2003 to correct PR field.)

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Sequence 34 BP; 0 A; 0 C; 2 G; 32 T; 0 U; 0 Other;

Gaps í, Length 34; IndelB Score 30.8; DB 1; Pred. No. 1.8e+02; 0; Mismatches 2; ô 1.1%; Conservative Local Similarity es 32; Conserv Query Match Best Loc Matches

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2706 ACTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2739 

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RESULT 85 AAD27124

AAD27124 standard; RNA; 37

AAD27124;

(first entry) 09-APR-2002 RNA template, (AU)3 used to direct RNA synthesis by HCV RNA polymerase.

HCV replicase; non-structural protein 5B; NS5B; polymerase; 88. Hepatitis C virus; lead compound; RNA

Unidentified

US6322966-B1

27-NOV-2001.

99US-00309670. 11-MAY-1999;

99US-00309670 11-MAY-1999; 

ZHONG W. (ZHON/) Z (HONG/) H (LAUJ/) I

HONG Z. LAU J Y N.

Lau JYN; Hong Z, Zhong W,

WPI; 2002-096587/13.

Assay system for hepatitis C virus replicase activity comprises RNA template with unstable, small stemloop capable of forming copy-back structure, viral non-structural protein 5B, nucleoside triphosphates. buffer.

Example 1; Fig 2A; 10pp; English.

The present invention relates to an assay system for hepatitis C virus C (HCV) replicase activity. The assay system comprises an RNA template that has an unstable, small stemloop at the 3 ' end capable of forming a copylack structure, a HCV non-structural protein 5B (NS5B), ATP, GTP, GTP, C and UTP nucleoside triphosphates (NTPS), where one of the NTP is activity of radiolabelled and an assay buffer that supports replication activity of NS5B. The invention also relates to the identification of optimal properties of an RNA template for copy-back self-priming RNA synthesis of HCV. This activity can be used to screen for anti-HCV replicase compounds. The C representation of comparates can be used for developing a system to characterise the biological relevance of lead compounds. The C representation of the invention is useful for developing a system to characterise the invention is useful for detecting HCV replicase activity. Semila RNA molecules to co-crystallise with HCV NS5B polymerase. The assay system of the invention is useful for detecting HCV replicase activity. The nucleic acid synthesised by NS5B is detected by evaluating an

Sequence 31 BP; 9 A; 9 C; 6 G; 6 T; 0 U; 1 Other;

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The present invention describes a nucleic acid segment of 10-100 contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a polymorphic site or an immediately adjacent base, or the complement of the segment. Also described are: (1) an allelesspecific oligonucleotide that hybridises to a segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the reference base indicated in the specification; and (3) analysing a nucleic acid, comprising obtaining a nucleic acid from an individual, and determining a base occupying any one of the polymorphic sites of the novelty. The nucleic acid segments and method can be used to analyse an individuals nucleic acid segments and method can be used to test for a disease phenotype and correlate the correlation of the phenotype with a particular polymorphism. The presence of polymorphic sites are useful for, e.g. forensics, paternity testing, correlation of phenotypic traits, AAA78631 to AAA79262 represent sequence correlation of phenotypic traits. AAA78631 to AAA79262 represent sequence of base occupying the polymorphic sites indicated using IUPAC-IUB
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autoradiograph of reaction products separated by gel electrophoresis. The present sequence is RNA template, (AU)3 used to direct RNA synthesis by RNA polymerase proteins of HCV, BVDV and polivirus. This sequence is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; genomic DNA; polymorphism; genome; allele-specific; primar; probe;
hybridisation; polymorphic site; forensic; paternity testing; medicine;
phenotypic trait; genetic analysis; genetic mapping; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genomic polymorphic nucleic acid segments, allele specific primers and probes, and methods of analysis, useful for e.g. forensics, paternity testing, genetic mapping,.
                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human genomic DNA polymorphic site sequence tag SEQ ID NO:566.
                                                                                                                                                Query Match 1.1%; Score 30.8; DB 1; Length 37; Best Local Similarity 94.1%; Pred. No. 1.9e+02; Matches 32; Conservative 0; Mismatches 2; Indels
                                                                                                         Sequence 37 BP; 32 A; 0 C; 2 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                       in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                             AAA79196 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AFFY-) AFFYMETRIX INC.
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The present invention describes a nucleic acid segment of 10-100 contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a polymorphic site or an immediately adjacent base, or the complement of the segment. Also described are: (1) an allelerspecific oligonucleotide that hybridises to a segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the reference base indicated in the specification; and (3) analysing a cucleic acid, comprising obtaining a nucleic acid from an individual, and determining a base occupying any one of the polymorphic sites of the covelty. The nucleic acid segments and method can be used to analyse an individuals nucleic acid segments and method can be used to test for a disease phenotype and correlate the presence of the phenotype with a particular polymorphism. The presence of polymorphisms with phenotypic traits and for genetic mapping of phenotypic traits. AAA78631 to AAA79262 represent sequence tags of human genomic DNA fragments containing polymorphic sites. The presence of buse occupying the polymorphic site is indicated using IUPAC-IUB
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hybridisation, polymorphic site, forensic, paternity testing, medicine,
phenotypic trait, genetic analysis, genetic mapping, ds.
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                                          Indels
     Length
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 Score 30.6; DB 1;
Pred. No. 1.8e+02;
                                                                          1716 TACTGCTGAAGAACACCATTACCTGAGGCC 1746
                                                                                               1 TACTGCTGAAGAAACRCCATTACCTGAGGCC 31
                                      1; Mismatches
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96.8%;
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                                    30; Conservative
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Best Local Similarity 96.8
Matches 30; Conservative
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Query Match
Best Local Similarity
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                                        Matches
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where the segment comprises a polymorphic site or an immediately adjacent
base, or the complement of the segment. Also described are: (1) an allele
capecific objigonucleotide that hybridises to a segment of the novelty.

capecific objigonucleotide that hybridises to a segment of the novelty.

capecific objigonucleotide that hybridises of the novelty where
capecific objigonucleotide that hybridises as sequence of the novelty where
capecific objigonucleotide that hybridises and by a base other than
the sequence is occupied by a base other than
the specification; and (3) analysing a
nucleic acid, comprising obtaining a mucleic acid from an individual, and
capecific occupying any one of the polymorphic sites of the
coverlay. The nucleic acid sequences for the presence of polymorphic sites of the
coverlay. The nucleic acid sequences for the presence of polymorphisms. The
coverlay of the phenotype with a particular polymorphism. The presence of
polymorphic sites are useful for, e.g. forensics, paternity testing,
correlation of polymorphisms with phenotypic ratals and for genetic
apping of phenotypic traits. AAA78631 to AAA7262 represent sequence
tags of human genomic DNA fragments containing polymorphic sites. The
                                                                                                                                                                                                                                                                                                           Human; genomic DNA; polymorphism; genome; allele-specific; primer; probe; hybridisation; polymorphic site; forensic; paternity testing; medicine; phenotypic trait; genetic analysis; genetic mapping; ds.
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                                                                                                                                                                                                                                                                        Human genomic DNA polymorphic site sequence tag SEQ ID NO:565.
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2201 AAAAGACTGGGCTCCTTGGTGGATGAGTTTA 2231
                     1 AAAAGACTGGGCTCCYTGGTGGATGAGTTTA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 21; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shah N, Warrigton JA;
                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JAN-2000; 2000EP-00250023.
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                                                                                                                                             AAA79195 standard; DNA; 31
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                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                         AAA79195;
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contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a nucreate or an immediately adjacent where the segment comprises a polymorphic site or an immediately adjacent base, or the complement of the segment. Also described are: (1) an allele capecific oligonuclectide that hybridises to a segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the reference base indicated in the specification; and (3) analysing a nucleic acid, comprising obtaining a nucleic acid from an individual, and determining a base occupying any one of the polymorphic sites of the novelty. The nucleic acid sequences for the presence of polymorphisms. The conventy and the sequence for the presence of polymorphisms. The presence of the phenotype with a particular polymorphism. The presence of the phenotype with a particular polymorphism. The presence of polymorphisms with a particular polymorphism. The presence of the phenotypic traits. AAA78631 to AAA79262 represent sequence of human senomic sequence of thusen secondariance of presence of human senomic sequence of the phenotypic traits. AAA78631 to AAA79262 represent sequence of human secondariance of the presence of human secondariance of the presence of presence of presence of presence of polymorphisms with phenotypic traits and for genetic traits the presence of human secondariance of the presence of polymorphisms with phenotypic traits and for genetic traits the presence of presence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human genomic polymorphic nucleic acid segments, allele specific primers and probes, and methods of analysis, useful for e.g. forensics, paternity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention describes a nucleic acid segment of 10-100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 21; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Warrigton JA;
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AAA79199 standard; DNA; 31 BP.
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                                                                                                                                                                    (first entry)
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                                                                                      AAA79199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patil N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Matches
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ID AAA7
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Gaps

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CCTAGCGATACCAAGAGGTCTCAGATCTATG 1622

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Query Match Best Local Similarity 96.8 Matches 30; Conservative

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<u>:</u>

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Human, genomic DNA, polymorphism, genome, allele-specific, primer, prob
hybridisation, polymorphic site, forensic, paternity testing, medicine,
phenotypic trait, genetic analysis, genetic mapping, ds.
                          Human genomic DNA polymorphic site sequence tag SEQ ID NO:563
                                                                                                                                                  Warrigton JA;
                                                                                                          26-JAN-2000; 2000EP-00250023.
                                                                                                                       99US-00238402
             (first entry)
                                                                                                                                                                                          testing, genetic mapping,.
                                                                                                                                    (AFFY-) AFFYMETRIX INC
                                                                                                                                                               WPI; 2000-500198/45.
                                                                                                                                                  Shah N,
             20-NOV-2000
                                                                                                                       27-JAN-1999;
                                                                   Homo sapiens
                                                                              EP1024200-A2
                                                                                             02-AUG-2000.
AAA79193;
                                                                                                                                                 Patil N,
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Human genomic polymorphic nucleic acid segments, allele specific primers and probes, and methods of analysis, useful for e.g. forensics, paternity Claim 1; Page 21; 141pp; English.

The present invention describes a nucleic acid segment of 10-100 contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a polymorphic site or an immediately adjacent base, or the complement of the segment. Also described are: (1) an allelectory of the complement of the segment. Also described are: (1) an allelectory of the complement of the segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the sepecification; and (3) analysing a nucleic acid, comprising obtaining a nucleic acid from an individual, and determining a base occupying any one of the polymorphic sites of the novelty. The nucleic acid segments and method can be used to analyse an individuals nucleic acid segments and method can be used to analyse and individuals nucleic acid segments for the presence of polymorphisms. The presence of the phenotype and correlate the presence of the phenotype in a particular polymorphism. The presence of polymorphisms with phenotypic traits and for genetic mapping of phenotypic traits and for genetic mapping of phenotypic traits. AAA78611 to AAA78252 represent sequence tags of human genomic DNA fragments containing polymorphic sites. The base occupying the polymorphic site is indicated using IUPAC-IUB

Gaps .; 0 1.1%; Score 30.6; DB 1; Length 31; 96.8%; Pred. No. 1.8e+02; Indels Sequence 31 BP; 5 A; 6 C; 8 G; 11 T; 0 U; 1 Other; ö Pred. No. 1.8e 1; Mismatches Local Similarity 96.8 Query Match Matches

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AAA79198 standard; DNA; 31 BP AAA79198; AAA79198 RESULT 田林和林林は田

(first entry) 20-NOV-2000

Human genomic DNA polymorphic site sequence tag SEQ ID NO:568.

Human, genomic DNA, polymorphism, genome, allele-specific, primer, probe, hybridisation, polymorphic site, forensic, paternity testing, medicine, phenotypic trait, genetic analysis, genetic mapping, ds.

Homo sapiens

EP1024200-A2

02-AUG-2000.

26-JAN-2000; 2000EP-00250023.

99US-00238402 27-JAN-1999;

(AFFY-) AFFYMETRIX INC.

Patil N, Shah N, Warrigton JA;

WPI; 2000-500198/45.

Human genomic polymorphic nucleic acid segments, allele specific primers and probes, and methods of analysis, useful for e.g. forensics, paternity

Claim 1; Page 21; 141pp; English.

testing, genetic mapping,.

The present invention describes a nucleic acid segment of 10-100 contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a polymorphic site or an immediately adjacent base, or the complement of the segment. Also described are: (1) an allelespecific oligonucleotide that hybridises to a segment of the novelty; (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the reference base indicated in the specification; and (3) analysing a nucleic acid, comprising obtaining a nucleic acid from an individual, and determining a base occupying any one of the polymorphic sites of the novelty. The nucleic acid sequences for the presence of polymorphisms. The method can also be used to test for a disease phenotype and correlate the presence of the phenotype with a particular polymorphism. The presence of polymorphisms with phenotypic traits and for genetic mapping of phenotypic traits. AAA78631 to AAA79262 represent sequence tags of human genomic DNA fragments containing polymorphic sites. The pass occupying the polymorphic site is indicated using IUPAC-IUB nomenclature 

Sequence 31 BP; 9 A; 7 C; 5 G; 9 T; 0 U; 1 Other;

ö 1.1%; Score 30.6; DB 1; Length 31; 96.8%; Pred. No. 1.8e+02; 0; Indels 1; Mismatches Conservative Local Similarity 30; Query Match Matches

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Gaps

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AAA79194 standard; DNA; 31 (first entry) 20-NOV-2000 AAA79194; RESULT 92 AAA79194 

ВР.

Human genomic DNA polymorphic site sequence tag SEQ ID NO:564.

Human, genomic DNA, polymorphism, genome, allele-specific, primer, probe, hybridisation, polymorphic site, forensic, paternity testing, medicine, phenotypic trait, genetic analysis, genetic mapping, ds.

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The present invention describes a nucleic acid segment of 10-100 contiguous bases chosen from one of 632 fragments (AAA78631 to AAA79262), where the segment comprises a polymorphic site or an immediately adjacent base, or the comprises a polymorphic site or an immediately adjacent especific oligonucleotide that hybridises to a segment of the novelty. (2) an isolated nucleic acid comprising a sequence of the novelty where the polymorphic site within the sequence is occupied by a base other than the reference base indicated in the specification; and (3) analysing a nucleic acid, comprising obtaining a nucleic acid sequences for the polymorphic sites of the novelty. The nucleic acid segments and method can be used to analyse an individuals nucleic acid segments and method can be used to analyse an individuals nucleic acid segments and method can be used to analyse and individuals nucleic acid sequences for the presence of polymorphism. The presence of the phenotype with a particular polymorphism. The presence of the phenotype with a particular polymorphism. The resence of polymorphic sites are useful for, e.g. forensics, paternity testing, correlation of polymorphic traits. AAA79262 represent sequence tags of human genomic DNA fragments containing polymorphic reals. The case of human genomic DNA fragments containing polymorphic reals.
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                                                                                                                                                                                                                                                                                                  Human genomic polymorphic nucleic acid segments, allele specific primers and probes, and methods of analysis, useful for e.g. forensics, paternity testing, genetic mapping,.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; single nucleotide polymorphism; SNP; forensic; paternity testing; genetic mapping of phenotypic trait; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          occupying the polymorphic site is indicated using IUPAC-IUB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1457 GIGATCTCTGTGGCCATTTATAATCTGGTCC 1487
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                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 21; 141pp; English.
                                                                                                                                                                                                                              Warrigton JA;
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                                                                                                              26-JAN-2000; 2000EP-00250023
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Best Local Similarity 96.8
Matches 30; Conservative
                                                                                                                                                                                        (AFFY-) AFFYMETRIX INC
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                                                                                                                                                                                                                              Shah N,
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Homo sapiens.
                                  EP1024200-A2
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The invention describes a nucleic acid molecule comprising one of 525 31 nucleotide sequences, given in the specification, or at least 10 nucleotides in length, and comprising a polymorphic site, where the nucleotide at the polymorphic site is different from a nucleotide at the polymorphic steens the sequence allele. The nucleotide at the polymorphic steens are useful in forensics comprising a single nucleotide polymorphism are useful in forensics (e.g. to identify an individual), in paternity testing, in correlating polymorphisms with phenotypic traits, and in genetic mapping of phenotypic traits. The correlation between a particular polymorphic form of a gene and a phenotype can be used in the diagnosis of that phenotype, as well as in the development of treatments for the phenotype. This sequence represents a fragment of a human gene found to containing a single nucleotide polymorphism following re-sequencing. The regions can be used to develop primers and probes for use in detect the SNP regions
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                                                                                                                                                              New nucleic acids comprising single nucleotide polymorphisms, useful in forensics (e.g. to identify an individual), paternity testing, correlating polymorphisms with phenotypic traits, and genetic mapping of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; urine sample analysis; kidney disease; glomerulonephritis; nephrotic syndrome; diabetes; lupus; hypertension; beta-actin; acute tubular necrosis; renal cancer; PCR; primer; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 30.6; DB 1;
Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
                                                                (WHED ) WHITEHEAD INST BIOMEDICAL RES.
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                                                                                                 Lander ES;
                                                                                                                                                                                                                                                 Example; Page 10; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                 24-JUL-2000; 2000US-0220315P.
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24-JUL-2001; 2001US-00912263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACF04897 standard; DNA; 32
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Best Local Similarity 96.8°
Matches 30; Conservative
                                                                                               Ireland JS,
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                                                                                                                                WPI; 2003-492161/46.
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                                                                                                                                                                                                                    phenotypic traits.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in individuals
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                                                                                                 Cargill M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 94
ACF04897/c
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ŏ Detecting or diagnosing a kidney disease, e.g. renal cancer or glomerulomphirtis, comprises determining the presence of expression podocyte gene for nephrin or proximal tubular cell gene for Indian hedgehog in a urine sample.

Claim 39; Page 24; Opp; English.

The present invention relates to a method of detecting a kidney disease, which comprises screening a mammalian urine sample for expression of a specific gene that is present in the urine sample only when cells indicating kidney disease are present, where the concentration of detectable albumin in the urine sample has a range of 0-30 mg/dl. The method is useful for detecting or diagnosing a kidney disease or disorders associated with e.g. glomerulonephritis, nephritic syndrome, diabetes, lupus, hypertension, acute tubular necrosis, renal obstructive disorders, renal cancers, and other diseases or symptoms. The podocyte gene for nephrin or the proximal tubular cell gene for Indian hedgehog is useful as selectable markers for a kidney disease. The present sequence is a PCR primer used to detect the human beta-actin gene

Sequence 32 BP; 2 A; 0 C; 1 G; 29 T; 0 U; 0 Other;

Gaps ; 0 1.1%; Score 30.4; DB 1; Length 32; 96.9%; Pred. No. 1.8e+02; Indels 2707 CTPABABABABABABABABABABABABA 2738 96.9%; Preq. ... Query Match Best Local Similarity 96.9\* \*\*+ches 31; Conservative ð

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AAS17761 standard; DNA; 31 BP RESULT 95

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12-MAR-2002 (first entry) AAS17761; 

Oligo d(T) PCR primer.

Oligo d(T); ss; differential subtraction; PCR primer; double exponential elimination; tumour

Synthetic.

US6316192-B1

13-NOV-2001.

99US-00268505 11-MAR-1999; 99US-00268505

11-MAR-1999;

(LUOJ/) LUO J

Luo J;

WPI; 2002-074371/10.

Selective elimination of non-targeted DNA sequences for rapid isolation and enrichment of the differences of DNA fragments between two pools of DNA, comprises converting testers to drivers.

Claim 6; Col 5; 23pp; English

of DNA fragments between two pools of DNA, comprises converting undesirable testers (DNA being subtracted) to drivers (DNA used to subtracted) and re-utilising converted drivers in repeats of subtraction to achieve double exponential elimination of undesirable tester sequences. invention comprises rapid isolation and enrichment of the differences The method comprises (a) attaching a nucleic acid fragment to 1 or more

CC polymerase chain reaction (PCR) adapters to form an adapter-attached nucleic acid fragment, followed by amplifying the adapter-attached nucleic acid fragment through PCR with primers containing nucleic acid capter through PCR with primers containing nucleic acid capter through PCR with primers containing nucleic acid capter that contains no attached adapter that contains no attached adapter whose sequence differs from the adapter to form a nucleic acid mixture, (d) adding to the nucleic acid mixture, (c) denaturing and re-annealing the tester/drive nucleic acid mixture, (d) adding to the nucleic acid mixture an effective amount of reagents necessary for removing the adapter sequence from the tester/ driver hetero-duplex and (e) repeating of step (c) to (d) at least once (no amplification takes place and no additional driver is added). The method is used for rapid isolation and enrichment of the differences of DNA fragments between two pools of DNA contribution in the search for tumour specific sequences. The method has 2 improvements over the methods disclosed by Anny et al. (1993), Straus et al. (1990) by (i) bypassing the need of a colymerase chain reaction (PCR) amplification or physical separation of desirable testers from undesirable ones in each repeat of climinates the necessity of tester dilution in each repeat of subtraction, it eliminates the need of subtraction. The present of subtraction. The present of subtraction. The present of subtraction. The present of sequence is an Oligo D(T) PCR primer used in the method of the invention

Sequence 31 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 1 Other;

Gaps ö Score 30.2; DB 1; Length 31; Pred. No. 1.9e+02; 1; Mismatches 0; Indels 96.8%; 30; Conservative Local Similarity Query Match Matches

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ઠે 용 AEE64867

AEE64867 standard; DNA; 31

AEE64867;

09-FEB-2006 (first entry) 

cDNA first strand synthesis primer.

ss; primer; gene expression; diagnosis.

Synthetic.

WO2005118791-A1.

15-DEC-2005.

17-MAR-2005; 2005WO-JP004788.

03-JUN-2004; 2004JP-00165208.

(AGEN ) NAT INST RADIOLOGICAL SCI.

Abe M, Araki R;

WPI; 2006-047539/05.

Producing gene expression profile by high coverage expression profiling analysis, by increasing mRNA in sample by obtaining amplified RNA complementary to cDNA using RNA polymerase, increasing double-stranded cDNAs having primers by PCR.

Example; SEQ ID NO 1; 49pp; Japanese.

The invention relates to a method of producing a gene expression profile. The method is useful for producing gene expression profile. The method is

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useful in the diagnosis of pathological conditions. The method enables preparation of cellular gene expression profile from an extremely small number of cells, pathological samples, micro tissues, etc., whose handling has been infeasible because of limited sample amount. The gene-expression profile can be obtained from trace amounts of sample by improved high coverage gene-expression profile analysis method. The method has high detection sensitivity with respect to gene expression. The present sequence represents a cDNA first strand synthesis primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cod liver uracil-DNA glycosylase enzyme, useful in monitoring or
rolling a reaction system multiplying DNA sequences or in carry-over
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heat-labile uracil-DNA glycolsylase; UNG; UDG; PCR primer; SMART; PCR control; LCR control; ligase chain reaction; carry-over prevention;
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                                                                                                                                                                                         1.1%; Score 30.2; DB 1; Length 31;
llarity 96.8%; Pred. No. 1.9e+02;
Conservative 1; Mismatches 0; Indels
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                                                                                                                                                            Seguence 31 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                 2708 ТАВАВАВАВАВАВАВАВАВАВАВАВАВА
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                                                                                                                                                                                                                                                                                                    31 ВАААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Guddal PH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Page 20; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JAN-2001; 2001WO-NO000008.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-JAN-2000; 2000NO-0000163.
27-OCT-2000; 2000NO-00005428.
                                                                                                                                                                                                                                                                                                                                                                                        AAS09500 standard; DNA; 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Willasen NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevention procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-451854/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SMART PCR primer #2
                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOT-) BIOTEC ASA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200151623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 controlling a
                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-OCT-2001
                                                                                                                                                                                                                               30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                           AAS09500;
                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lanes O,
                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                        RESULT 97
                                                                                                                                                                                                                                                                                                                                                                        AAS09500,
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The present invention relates to fibrinolytic enzyme, brevinase (see AAG79000), which is separated from the poison of Agkistordon blomhoffi brevicaudus (mamushi). The enzyme shows stability at high temperatures and is thus useful in developing thrombolytic agents. The present sequence is a PCR primer, which was used in an example from the present invention. (Updated on 11-SEP-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                     Fibrinolytic enzyme; brevinase; thermostable; thrombolytic agent; mamushi; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                   Fibrinolytic enzyme, brevinase, separated from poison of viper,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 32;
                                                                                                brevinase, PCR primer, BbRP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IndelB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 32 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 30.2; DB 1;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2708 TAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence of scissile link probe MRC064 (HL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Agkistrodon blomhoffi; brevicaudus
                                                                                                                                                                                                                                                                                                                                                               agkistrodon blomhoffi brevicaudus.
                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 6; 23pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                           99KR-00049115
                                                                                                                                                                                                                                                  99KR-00049115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%;
96.8%;
                                                                                                Mamushi fibrinolytic enzyme,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1277/c
AAN70277 standard; DNA; 30
                   ABA01204 standard; DNA; 32
                                                                 (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybridisation; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (revised)
                                                                                                                                                                                                                                                                                                                              WPI; 2001-636862/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                          Park W;
                                                                                                                                                                                                                                                                       (LEEJ/) LEE J W.
(PARK/) PARK W.
                                                                                                                                                                               KR2001045716-A.
                                                                                                                                                                                                                           06-NOV-1999;
                                                                                                                                                                                                                                                  06-NOV-1999;
                                                                11-SEP-2003
28-JAN-2002
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                                          ABA01204;
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                                                                                                                                                                                                                                                                                                          Lee JW,
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RESULT 98
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Probe MRCO64 is bound by a hydrolysable linkage to a solid support at its 3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to from 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the libonucleotides by a ds RNase, eg RNase H or ExoIII. (Updated on 13-0CT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Overexpression of the enzyme glutathione-s-transferase pi has been implicated as being responsible for the broad range drug resistance which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic oligo-nucleotide(s), prodn. useful e.g. for HIV-1 inhibition which bind to target sequence in duplex DNA forming colinear triplex by binding to major groove.
                                                                                                                               Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Glutathione-s-transferase pi, cancer; drug resistance; chemotherapy; sensitisation; triplex; target; duplex; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 0 A; 0 C; 0 G; 22 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%; Score 30; DB
100.0%; Pred. No. 1.9
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GST3anti, for GSTpi gene target sequence.
                                                                                                                                                                                   Disclosure; Page 24; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Col 27; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89US-00453532
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ36302 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30; Conservative
                      MEIOGENICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kessler DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1993-035718/04.
                                                                                             WPI; 1989-339977/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                           Bender R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BAYU ) BAYLOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-DEC-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-DEC-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
07-JUN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hogan ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ36302;
                      (MEIO-)
                                                           Duck P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ36302,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                               The patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- = a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the samples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                   Synthetic nucleic acid probes - comprising two nucleic acid sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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0
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100.0%; Pred. No. 1.9e+02;
ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 0 A; 0 C; 0 G; 22 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag= a
note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                Robertson JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30 AAAAAAAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Probe MRCO64; solid support; ribonuclease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= b
/note= "ribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cocation/Qualifiers
                                                                                                                                               Crosby W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВР
                                                                                                                                                                                                                                      linked by a scissile linkage.
                                    86EP-00116906
                                                                        85US-00805279
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                                                                                                                                                                                                                                                                          Example; p29; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAN92243 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
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Matches 30; Conservative
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/*tag= c
                                                                                                        (MEIO-) MEIOGENICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                                                                                                 WPI; 1987-186567/27
                                                                                                                                             Bender R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS probe MRCO64.
                                  04-DEC-1986;
                                                                    05-DEC-1985;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-APR-1988;
08-JUL-1987
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25-APR-1990
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                                                                                                                                             Duck P,
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AAN92243

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Gaps

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DB 1; Length 30; 1.9e+02; 0; Indels

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develops in a variety of cancers. Expression of the gene may be prevented by the formation of a triplex between the duplex target DNA sequence and an anti parallel or parallel synthetic oligonucleotide. A suitable target sequence is that from base -499 to -410 of GSTpi, an unusual repetitive DNA segment within the control domain. Oligonucleotides targetted against this sequence will repress GSTpi transcription. See also AAQ36219-362. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic oligo-nucleotide(s), prodn. useful e.g. for HIV-1 inhibition .
which bind to target sequence in duplex DNA forming colinear triplex by
                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glutathione-s-transferase pi, cancer; drug resistance; chemotherapy; sensitisation; triplex; target; duplex; ss.
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                                                                                                                                                                             DB 1; Length 30;
1.9e+02;
                                                                                                                                                                                                             Indels
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                                                                                                                                            Seguence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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                                                                                                                                                                                       100.0%; Prea. ...
                                                                                                                                                                                                                                                                     GST3par, for GSTpi gene target sequence.
                                                                                                                                                                              1.1%; Score 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Col 27; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BAYU ) BAYLOR COLLEGE MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         89US-00453532
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                                                                                                                                                                                                                                                                                                                                                                        AAQ36301 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   binding to major groove.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hogan ME, Kessler DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1993-035718/04.
                                                                                                                                                                                               Local Similarity
les 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-DEC-1989;
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07-JUN-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                  30
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                                                                                                                                                                                                               Matches
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Vaccine, cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; tumous fungal infection; parasitic infection; cancer; asthma; infections disease; allergy; immune deficiency; phosphorothioate; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                      Optical detection of hybridization complexes for specific target nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target nucleic acid presence in a sample. A preferred target is a Mycobacterium complex nucleic acid sequence. The detection method uses visual detection of a change in the hybridization without aid of instrumentation. Multiple copies of a target nucleic acid sequence are mixed with first and second detectable probes under hybridizing conditions favouring particle agglutination via a bridging molecule allowing for visual detection of the target nucleic acid sequence. The bridging molecule enhances or inhibits formation of a hybridization
                                                                                                                                          Visual, nucleic acid detection; target; hybridisation; probe; primer; agglutination; bridging molecule; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel method for the visual detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 30; DB 1;
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                                                                                                                                                                                                                                                                                                                                                                   Reshatoff MR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunostimulatory nucleic acid #1005.
                                                                                                                WO9923258 oligonucleotide primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 40; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                             98WO-US023267.
                                                                                                                                                                                                                                                                                                          97US-0063969P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 30
                            AAX57020 standard; DNA; 30
                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30; Conservative
                                                                                                                                                                                                                                                                                                                                                                   Stull PD,
                                                                                                                                                                                                                                                                                                                                      (GENP-) GEN-PROBE INC.
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-326994/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF99889 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            acid sequences
                                                                                                                                                                                                                                                                                                                                                                    Weisburg WG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-JUN-2001
                                                                                                                                                                                                                                                                                                          31-OCT-1997;
                                                                                                                                                                                                                                                                             30-OCT-1998;
                                                                                     19-JUL-1999
                                                                                                                                                                                                                     WO9923258-A1
                                                                                                                                                                                                                                                14-MAY-1999.
                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF99889;
                                                         AAX57020;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      complex
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RESULT 10
AAX57020/
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Gaps

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Indels

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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738

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1.1%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 1.9e+02;

100.0%; Pred. No. 1.3 ive 0; Mismatches

Best Local Similarity Matches 30; Conserv

Query Match

05-APR-2001

Synthetic.

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response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immuno response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. berpesviridae, retroviridae haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                             Vaccinating against tumors, infectious diseases, allergies and asthmausing immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for stimulating an immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining an analyte in a sample, for generating multiple double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88; 5'-A30-3'; double stranded DNA generation; promiscuous base;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 1.9e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic primer sequence 5'-A30-3'.
                                                                                                                                                                                                                                                                                                                                                    Example 6; Page 60; 338pp; English
                                                                                                                                                                                      Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HOFF ) ROCHE DIAGNOSTICS GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВР
                     25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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                                                                                                                  IOWA RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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Matches 30; Conservative
                                                                                                             (IOWA ) UNIV IOWA RES FO
(COLE-) COLEY PHARM GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         target molecule; primer
                                                                                                                                                                                      Schetter C,
                                                                                                                                                                                                                                    WPI; 2001-273485/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-214947/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6326143-B1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-DEC-2001.
                                                                                                                                                                                      Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Orum H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acid. The immunostimulatory nucleic acid can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. toxoplasma, namoprilus, campylobacter, clostridium, Bscherichia coli and/or staphylococus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccinating against tumors, infectious diseases, allergies and asthmausing immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30 BP; 30 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. .v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 AAAAAAAAAAAAAAAAAAAAAAAAAAA 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunostimulatory nucleic acid #1004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 60; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                       Vollmer J;
                                                                                                                                                                                                    25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF99888 standard; DNA; 30 BP
                                                                                                                                                         25-SEP-2000; 2000WO-US026383.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-2000; 2000WO-US026383.
                                                                                                                                                                                                                                                                                              (IOWA ) UNIV IOWA RES FOUND.
(COLE-) COLEY PHARM GMBH.
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                                                                                                                                                                                                                                                                                                                                                                     Krieg AM, Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-273485/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 30; Conserv
                                                               WO200122972-A2
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12-JUN-2001

AAF99888;

Query Match

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Gaps

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Tue Nov

stranded nucleic acids, comprises employing a single primer sequence with a nucleobase sequence having affinity to the sequence contained in a target nucleic acid.

Example 1; Col 14; 25pp; English

The intention relates to determining an analyte in a sample computation of the providing a target nucleic acid comprising a region A, a nucleobase sequence B, and a sequence I linked to the 5' terminus of the nucleobase sequence B, where the nucleobase sequence B is not specific for the sequence B, where the nucleobase sequence B is not specific for the sequence B, where the sample, (d) binding the target nucleic acid from the remaining part of the sample, (d) the target nucleic acid from the remaining part of the sample, (d) the formal a primer to the nucleobase sequence B', and the nucleobase sequence B', and the nucleobase sequence B' and the nucleobase sequence B', and the nucleobase sequence B' as a template and using nucleotides, where at least 30 % of the nucleotides contain at least one promiscuous base which is capable of searching with each of adenine, guanine, cytosine, and thymine, (E) separating the target nucleic acid from the elongation product E, where the elongation product E, where the elongation product E is capable of acting as a template for the elongation of the further primer, (h) separating the hybridised further primer of step (d) to produce an elongation product E, where the elongation product E is capable of acting as a template for the elongation product E, where at least 10 % of the nucleobase sequence B' (i) separating the elongation product E' (j) hybridising a template and using the elongation product E' (j) hybridising a template and using nucleotides, where at least 30 % of the nucleobase sequence B' (i) separating the elongation product E' (j) to produce C another elongation product E', (j) hybridising a further primer of step (j) to product E (m) product E as a template and using nucleotides, where at least 30 % of the nucleobase sequence B' (k) elongating the further primer of step (j) to product E, (m) could be a template and using nucleotides, where at least 30 % of the nucleobase sequence B' (j) to product E as a template and using nucleotides contain at least nucleobase determining the elongation product B and/or elongation product B' as a measure of the presence or amount of the analyte, where the lengths of the sequence I and the mucleobase sequence B are chosen such that, when the further primer bypridises to the elongation product E in step (g), the further primer spans a sequence formed by elongation of the Pybridised primer of step (e) and overlaps at least a part of the 3' region of the hybridized primer of step (e) by an overlap length. The method is useful determining an analyte in a sample. In particular, the method is useful for generating multiple double stranded nucleic acids. The present sequence is a primer molecule used to exemplify the method of the invention invention relates to determining an analyte in a sample comprising

Sequence 30 BP; 30 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö DB 1; Length 30; Indels 1.9e+02; 1.1%; Score 30; DB 100.0%; Pred. No. 1.9 :ive 0; Mismatches Juery Match
Best Local Similarity 100...
Best Anna 30; Conservative

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8 셤 RESULT 107 ABK10412/c

ABK10412 standard; DNA; 30 BP

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ABK10412;

21-MAY-2002 (first entry)

Synthetic primer sequence 5'-T30-3'. XZZZZZZZZX

ss; 5'-T30-3'; double stranded DNA generation; promiscuous base; target molecule; primer.

the invention trained to determining an analyte in a sumpte Comprising sequence B, and a sequence I linked to the 5' terminus of the nucleobase sequence B, and a sequence I linked to the 5' terminus of the nucleobase sequence B, and the region A specifically binds to the analyte, (b) binding the target nucleic acid from the remaining part of the sample, (d) binding the target nucleic acid from the remaining part of the sample, (d) the target nucleic acid from the remaining part of the sample, (d) the target nucleic acid from the remaining part of the sample, (d) the target nucleic acid from the remaining part of the sample. (d) the target nucleic acid from the remaining part of the sample. (d) the target nucleobase sequence B', and the nucleobase sequence B' the bingating the bybridised corrected on elongation product E using the target nucleic acid from the least 30 % of the nucleotides contain at least one promiscous base which is capable of sate template and using nucleotides of from the elongation product E; (g) separating the target nucleic acid from the elongation product E; where the elongation product E; (d) to produce an elongation product E; where the elongation product E is capable of acting as a template for the elongation of the further primer. (h) elongating the hybridised further primer of step (g) to produce and elongating the product E; (s) hybridising a further primer comprising a contain at least one promiscuous base, (i) separating the elongation product E; (k) hybridising a further primer comprising a nucleobase sequence B; to the target nucleic acid or the elongation product E; (k) elongating the further primer of step (j) to product C; product E as a template and using nucleotides, where at least 30 % of the nucleotides contain at least one promiscuous base, (l) a sufficient nucleotides contain at least one promiscuous base, (l) a sufficient number of teast one promiscuous (n) - (l) a sufficient number of teast one promiscuous (n) - (l) a sufficient number of teast one promiscuous (n) - ( Determining an analyte in a sample, for generating multiple double stranded nucleic acids, comprises employing a single primer sequence with a nucleobase sequence having affinity to the sequence contained in a optionally repeating steps (9) - (1) a sufficient number of times to optionally repeating steps (9) - (1) a sufficient number of times to generate a desired amount of double stranded nucleic acids and (n) determining the elongation product E and/or elongation product E' as a measure of the presence or amount of the analyte, where the lengths of the sequence I and the nucleobase sequence B are chosen such that, when the further primer hybridises to the elongation product E in step (g), the further primer spans a sequence formed by elongation of the hybridised primer of step (e) and overlaps at least a part of the 3, region of the hybridized primer of step (e) by an overlap length. The method is useful determining an analyte in a sample. In particular, the method is useful for generating multiple double stranded nucleic acids. The invention relates to determining an analyte in a sample comprising Example 1; Col 14; 25pp; English. (HOFF ) ROCHE DIAGNOSTICS GMBH 98US-00083123. 96WO-EP005149. WPI; 2002-214947/27. target nucleic acid. Seeger C; US6326143-B1 22-MAY-1998; 22-NOV-1996; 04-DEC-2001. Synthetic. Orum H, 

Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;

The present sequence is a primer molecule used to exemplify the method of the invention

method is useful determining an analyte in a sample. In method is useful for generating multiple double stranded

DB 1; Length 30; Score 30; DB 1; I Pred, No. 1.9e+02; Query Match Best Local Similarity 100.0 Matches 30, Conservative

Indels 100.0%; Prec. ...

2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738

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Gaps ö

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Fluorescent intercalative dye, nucleic acid detection, gene diagnosis, clinical diagnostics; Stokes shift; ds.

Method of measuring nucleic acid related oligonucleotide dT30mer.

(first entry)

26-NOV-2002

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comprising the sequence (CTATTTT) n and its complement (AAAATAG) n, where n is 1. The probe sets of the invention are useful for detecting kappa or lambda-immunoglobulin light chain mRNA or corresponding heteronuclear RNA, CWY (cytomegalovirus) immediate early RNA, EBW (Epstein-Barr virus) early RNA 1 and RNA 2, and human Alu repetitive satellite genomic sequences. The invention is a useful generic sequence for incorporation into oligonucleotide probes for detecting gene-specific sequences within cells or tissue samples in in airu hybridisation analysis and for attaching a label to immunoblatochemical analyses. The present nucleic acid sequence represents one of a collection (ABK70376-ABK70501) of oligonucleotide probes that were used in the invention for detecting or collisonucleotide probes that were used in the invention for detecting or localising a plurality nucleic acid target gene or antigen within a cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel oligonucleotide label-domain for incorporation into oligonucleotide probes useful for detecting or localizing nucleic acid target genes within a cell or tissue sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a new oligonucleotide label-domain
                                                                                                                                                                                                                                                     Human, oligonucleotide label-domain; CMV; cytomegalovirus; EBV,
Epstein-Barr virus; lambda-immunoglobulin light chain; hapten;
kappa-immunoglobulin light chain; repetitive Alu sequence; EBER;
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Pred. No. 1.9e+02;
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Matches 30; Conservative 0; Mismatches
    AAAAAAAAAAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                In-situ analysis synthetic probe #58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (VENT-) VENTANA MEDICAL SYSTEMS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 69; 71pp; English.
                                                                                                                                                                                                                                                                                                                  Epstein-Barr early RNA; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jtermohlen JG, Connaughton J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-SEP-2001; 2001WO-US028014.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-SEP-2000; 2000US-0233177P.
                                                                                                ABK70490 standard; DNA; 30
                                                                                                                                                                           15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-371972/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue sample
                                                                                                                                                                                                                                                                                                                                                                                              WO200222874-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                       21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                         Synthetic.
30
                                                                                                                                    ABK70490;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                         RESULT 108
                                                                             ABK70490
g
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The invention describes a novel fluorescent dye and method of detecting nucleic acid. The dye and method are useful for nucleic acid probes for measuring nucleic acid s) containing a specific nucleic acid probes for a measuring nucleic acid solution to a sample, and for qualitative/quantitative assay of target RNA containing a specific base sequence anticipated in gene mixture. The assay is useful contained diagnostics and in send diagnostics and in cidentification/quantification microoxganisms in biological samples such as serum, plasma and urine, microbially contaminated samples such as serum, plasma and urine, microbially contaminated samples such as serum, plasma and sea. The fluorescent intercalation dye shows a large fluorescent enhancement upon intercalation into double-stranded concleic acid, and shows a great difference between excitation and concleic acid, and shows a large Stokes shift) and does not have a fluorescent spectrum that overlaps with those of conventionally known cluorescent spectrum that overlaps with those of conventionally known cluorescent intercalation dyes. Viruses, microbial RNAs, specific sequences in one RNA, are detected or quantified in a short time, hence the detection method is applicable to clinical diagnosis which requires high reliability. Amplification and extraction efficiencies of the target on uncleic acid, are checked. This sequence represents a synthetic DNA used construction by a construction by an activity by a construction and extraction efficiencies of the target on a checked. This sequence represents a synthetic DNA used

Fluorescent dye or its salt, hydrate, solvate or stereoisomer for nucleic acid probe for measuring nucleic acid(s) containing specific nucleic acid sequence in sample, has specific formula.

Example 5; Page 33; 40pp; English

뭐;

Tokunaga T, Ishiguro T,

(TOYJ ) TOSOH CORP.

WPI; 2002-645688/70.

11-JAN-2002; 2002EP-00000723. 11-JAN-2001; 2001JP-00003432.

EP1223226-A2

Synthetic.

17-JUL-2002.

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3'-amino oligonucleotide; solid support; benzene derivative; solid phase oligonucleotide synthesis; SPOS; controlled pore glass; CPG;
                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                            Length 30;
                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide #2 synthesised on a solid support.
                                                                                                                                                                                                                                                                                                                              Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                           1.1%; Score 30; DB 1; Los 100.0%; Pred. No. 1.9e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                     2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                 30 AAAAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                 ADU07154 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                        30; Conservative
                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           27-JAN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                               ADU07154;
                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 110
                                                                                                                                                                                                                                                                                                                                                        Matches
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셤
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Gaps

2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738

8 셤

30 AAAAAAAAAAAAAAAAAAAAAAAAA 1

ABS53961 standard; DNA; 30 BP

RESULT 109 ABS53961/c

ABS53961;

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/mod_base= OTHER
/note= "OTHER= Linked to gold-S' where S' indicates a
connecting unit prepared via an epiandrosterone disulfide
                                                                                                                                                                                                                        The invention describes a microarray comprising QC probes and a method for fabricating the microarray. The microarray comprising QC probes has the complementary nucleotide sequence with that of a target gene or any nucleotide sequence material albeled to the target gene or any different from the fluorescence material labeled to the target gene, wherein the fluorescence material is labeled at the 3-terminal 55-terminal or intermediate of the QC probes, a spacer is further contained between the nucleotide sequence of the probes and fluorescence material; the QC probes can be cDNA, oligonucleotides, peptides or proteins. The method for fabricating the microarray comprising QC probes involves fixing the QC probes and target probe involves substrate, wherein the QC probes and target probe on a substrate, wherein the QC probes and target probe on a substrate, wherein the QC probes and target probe on a substrate, wherein the QC probes and target probe are fixed in a spot at the same time. This sequence represents an oligonucleotide associated with the microaray of the invariance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analyte detection; DNA detection; protein detection; MecA gene; probe;
                                                                                                                                                Microarray comprising QC probes and method for fabricating the same.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylococcus aureus MecA gene specific probe 1 SEQ ID:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Le
. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%; Scc...
100.0%; Pred. No. 1...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         АВАВАВАВАВАВАВАВАВАВАВАВАВ 1
                                                                                                                                                                                             Example 2; SEQ ID NO 9; 17pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the same time. This sequence represe:
with the microaray of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAY-2003; 2003US-0474569P.
29-AUG-2003; 2003US-0499034P.
04-NOV-2003; 2003US-0517450P.
23-MAY-2004; 2004US-0567874P.
27-MAY-2004; 2004US-00854848.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-2004; 2004US-00995051.
                                                               Park HK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AED67969 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.
nes 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1. .30
/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      group"
                                                             Jang HJ, Kim CM,
(KIMC/) KIM C M.
(PARK/) PARK H K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2005250094-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to the preparation of 3'-amino oligonucleotide derivatives. The method comprises providing solid support benzene derivatives. The method comprises providing solid support benzene derivatives, synthesising an oligonucleotide pursuant to standard techniques for solid phase oligonucleotide synthesis (SPOS) where the oligonucleotide chain is assembled on the solid support, cleaving the oligonucleotide from the solid support and deprotecting the oligonucleotide completely except for the terminal protective group. The oligonucleotides are useful in the synthesis of bi-fluorescent probes. This process suppresses the undealred formation of unmodified 3'-OH oligonucleotides through cyclic phosphate intermediates and reduces the probability of errors resulting from the use of different sets of oligonucleotides. This sequence represents an invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                           Preparation of 3'-amino oligonucleotide derivatives, useful to synthesis bi-fluorescent probes, comprises providing solid support compounds, synthesis an oligonucleotide chain is assembled on the solid support,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Microarray associated oligonucleotide SEQ ID NO 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%; Scc...
100.0%; Pred. No. -
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 АААААААААААААААААААААААААААА 2738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microarray; DNA detection; hybridization;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      арарарарарарарарарарарарарара 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADV98265 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-FEB-2004; 2004KR-00007237.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-FEB-2003; 2003KR-00006722.
                                                                                                                                                                             21-APR-2004; 2004US-00830484
                                                                                                                                                                                                                    21-APR-2003; 2003US-0464269P
                                                                                                                                                                                                                                                                                                                                                                                                                                                               cleavage and deprotection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
      bi-fluorescent probe; ss
                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-794500/78
                                                                                                                                                                                                                                                                   (PROL-) PROLIGO LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                              Wolter A;
                                                                                          US2004220397-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KR2004076201-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-AUG-2004
                                                                                                                                   04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                 Synthetic
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RESULT 111

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Gaps

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0; Indels

Length 30;

25 #2

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Storhoff
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The present invention provides a method for detecting the presence or absence of a single target molecule or target analyte (e.g. nucleic acid, protein, lipid, bacterium). The method involves contacting sample with one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is a Staphylococcus aureus MecA gene specific probe. This sequence is used in the preparation of nanoparticle-oligonacleotide conjugate probes. Note: The present sequence is the SEQ ID No: 25 shown on page 25 in example 25 of the specification. This sequence differs from the SEQ ID NO: 25 given in the sequence listing (see AED67954). Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light scattering complex formed. Garimella V; Σ Senical Bao YP, Example 25; SEQ ID NO 25; 70pp; English. Muller UR, Lucas A, WPI; 2005-784662/80.

.. 1.1%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 1.9e+02; ive 0; Mismatches 0; Indels Sequence 30 BP; 30 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 30; Conservative Similarity Query Match Best Local Matches ò

2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738 

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AED67958 standard; DNA; 30 12-JAN-2006 (first entry) AED67958; 

Analyte detection; DNA detection; protein detection; MecA gene; probe; Methicillin resistant S. aureus MecA gene specific probe 1 SEQ ID:

Staphylococcus aureus US2005250094-A1.

10-NOV-2005

22-NOV-2004; 2004US-00995051

30-MAY-2003; 2003US-0474569P. 29-NGC-2003; 2003US-0499034P. 04-NOV-2003; 2003US-0517450P. 03-MAY-2004; 2004US-0567874P. 27-MAY-2004; 2004US-00854848.

(NANO-) NANOSPHERE INC.

Garimella V; Senical M, Bao YP, Muller UR, Storhoff JJ, Lucas A, WPI; 2005-784662/80. Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light scattering complex formed.

Example 25; SEQ ID NO 29; 70pp; English.

The present invention provides a method for detecting the presence or absence of a single target molecule or target analyte (e.g. nucleic acid, protein, lipid, bacterium). The method involves contacting sample with one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is a methicillin resistant Staphylococcus aureus (MRSA) MecA gene specific probe. This sequence is used in the preparation of nanoparticle-oligonucleotide conjugate probes. Note: This sequence is incorrectly designated as SEQ ID NO: 25 in example 25 (page 25) of the specification 

Sequence 30 BP; 30 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Length 30; 0; Indels Query Match
1.1%; Score 30; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.9e+02;
Matches 30; Conservative 0; Mismatches 0;

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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738 

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839/c AEE86839 standard; DNA; 30 RESULT 114 AEE86839,

BP

(first entry) 23-FEB-2006 AEE86839;

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Gaps

Novel solid phase-related oligonucleotide Oligo dT40-Cy3 #14.

DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.

Synthetic.

Location/Qualifiers /note= "OTHER= Cy3" /\*tag= a /mod\_base= OTHER modified\_base

DE102004025746-A1. 15-DEC-2005. 

26-MAY-2004; 2004DE-10025746 (CHER/) CHERKASOV D.

(HENN/)

26-MAY-2004; 2004DE-10025746.

Cherkasov D, Hennig C, HENNIG C. GENOVOXX GMBH. (GENO-)

ü Baeuml

WPI; 2006-040183/05.

Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.

Disclosure; Page 97; 144pp; German.

This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged and repeated contact of the solid phase with high concentrations of labeled

88.

RESULT 115 **AEE86831**,

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This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged and respected contact of the solid phase with high concentrations of labeled reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parallel sequencing of nucleic acids by optical methods, by cyclic primer -marrix extension, using a solid phase with reduced non-specific binding of labeled components.
                                                                                                                                                                                                                                DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing;
                                                                                                                                                                                              Novel solid phase-related oligonucleotide Oligo dT40-Biotin #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB 1; Length 30;
Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= 5'-terminal Biotin-TEG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
0; Mismatches
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30 AAAAAAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 97; 144pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baeuml
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2004; 2004DE-10025746
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                                                                                       AEE86833 standard; DNA; 30
                                                                                                                                                                (first entry)
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Best Local Similarity 100.
Matches 30, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hennig C,
                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHER/) CHERKASOV D.
(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2006-040183/05
                                                                                                                                                                                                                                                                                                                                                                                                                  DE102004025746-A1
                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cherkasov D,
                                                                                                                                                                23-FEB-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-DEC-2005.
                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                               AEE86833
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                                                        RESULT 11
AEE86833/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged and respected contact of the solid phase with high concentrations of labeled reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.
   reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                       Novel solid phase-related oligonucleotide Oligo-dT30-Cy3 #6.
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100.0%; Pred. No. 1.9e+02;
tive 0; Mismatches 0; Indels
                                                                                           Length 30;
                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                        Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                           DB 1; Le
1.9e+02;
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100.0%; Pred. No. ...
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                              AEE86831 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                  Conservative
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nes 30, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HENNIG C.
GENOVOXX GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2006-040183/05
                                                                                       Query Match
Best Local Similarity
Matches 30; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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Matches
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Synthetic

(CHER/)

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This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonicleotide which was used in the development of the novel solid
                                                                                                                                                                                                                                                         Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel solid phase-related oligonucleotide Oligo-dT30-Cy3 #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 30; DB 1; L
Pred. No. 1.9e+02;
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100.0%; Pred. No. 1...
0; Mismatches
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/mod_base= OTHER
/note= "OTHER= Cy3"
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                                                      26-MAY-2004; 2004DE-10025745
                                                                                        26-MAY-2004; 2004DE-10025745
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEE86847 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  phase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hennig C;
                                                                                                                    (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
(CHER/) CHERKASOV D.
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(GENO-) GENOVOXX GMBH.
(CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                           WPI; 2006-040182/05
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid phase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components.
                               DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
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Novel solid phase-related oligonucleotide Oligo dT40-Biotin #7.
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                                                                                                                                                                     /note= "OTHER= 5'-terminal Biotin-TEG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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1.9e+02;
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v 100.0%; Pred. No. 1..
0; Mismatches
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                                                                                                   Location/Qualifiers
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                                                                                                                                        *tag= a
mod base= OTHER
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/mod_base= OTHER
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ID AEE86855 standard; DNA; 30
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                                                                                                                                      *tag=
                                                                                                                                                                                                                                                                                                                                      (HENN/) HENNIG C. (GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                       CHERKASOV D.
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Best Local Similarity
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modified_base
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modified_base
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Synthetic

AEE86855

RESULT 118

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Length 30; 0; Indels

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26-MAY-2004; 2004DE-10025695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel solid support surface for parallel analysis of many individual nucleic acids by optical methods. The invention also describes, a) a solid phase in which the surface shows reduced non-specific binding of labeled components; b) methods for preparing the novel solid support and c) methods of parallel analysis of
                                                                       This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced nor-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid
                    Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Surface of a solid support, useful for multiple parallel analysis of nucleic acids by optical methods, having low non-specific binding of labeled components.
                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "optionally labeled with Cy3 or Biotin-TEG"
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                                                                                                                                                                               DB 1; Length 30;
1.9e+02;
                                                                                                                                                                                                   Indels
                                                                                                                                                          Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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                                                                                                                                                                               Query Match
1.1%; Score 30; DB
Best Local Similarity 100.0%; Pred. No. 1.9
Matches 30; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                     DNA detection; DNA sequencing; primer; 8S.
                                                                                                                                                                                                                                            30 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
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                                                       Disclosure; Page 62; 88pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
                                                                                                                                                                                                                                                                                                  AEF12156 standard; DNA; 30 BP
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                                                                                                                                                                                                                                                                                                                                           09-MAR-2006 (first entry)
                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide dT30-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hennig C;
                                                                                                                                        phase of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHER/) CHERKASOV D.
    WPI; 2006-040182/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DE102004025744-A1.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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(GENO-)
                                                                                                                                                                                                                                                                                RESULT 120
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contracted acid by optical methods, using the solid support. The surface of the solid support is made of silica, glass, silicon dioxide or si-Odi; is flat and has uncleic acid chains fixed to it, optionally chrough a linker. The solid phase is preferably part of a device that allows fluid exchange and it is permeable to light in the wavelength cregions 200-400; 200-2000 or 400-800 min. An external layer of solid cupport is removed, then the nucleic acid is coupled to it, optionally cupport is removed, then the nucleic acid is coupled to it, optionally after attachment of a linker layer. Alternatively, after removing the external layer, nucleic acids are synthesized on the surface by cyclic coupling, optionally after attachment of a linker, and in either case, actachment or synthesis of nucleic acide on the surface layer attachment or synthesis of nucleic acide on the surface is not attachment or synthesis of nucleic acide on the surface is not direct containing monomers or polymers) can be compaid, especially to remove a layer 1 nm to 100 micron thick. Solutionally after removal of the surface layer, the surface is not direct and all subsequent steps are done in a liquid phase. The nucleic acids and lyzed represent a single population or many different populations and contains s-50, 20-200 or 50-500 nucleotides. The linker is 1-50 nm long can alize brinker-matrix complexes. The linker has a branched or linear polymer: (etrept) avidin or a nucleic acid etriphosphates, in which the label is cleavable. Particularly analysis involves cyclic sequencing and a preferred method contains 30-300 or t, RNA or DNA, and the solid support, with formation of a ceromatrively the nucleic acid is fared to the support and then actions and thermatrial sequences that function as primers for the sequencing and the achieve that primer. The innocrporated mucleotide can be contains 30-300 or t, RNA or DNA, and the solid by acception or alternatively the nucleic acid is fared to the support and then achieve the function of a reversible ss; dna detection; DNA sequencing; DNA amplification; oligo dT40-biot. Gaps Optical DNA analysis process-related oligonucleotide dT40-biotin. so assay sensitivity is improved. This sequence represents an oligonuclectide used to illustrate the method of the invention. ö Length 30; Indela Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other; . 1.1%; Score 30; DB 1; L 100.0%; Pred. No. 1.9e+02; 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738 100.0%; Prec. ... /\*tag= b /mod\_base= 5'-Biotin-TEG Location/Qualifiers BP. AEF94776 standard; DNA; 30 (first entry) 30; Conservative Best Local Similarity DE102004025695-A1 modified\_base Unidentified 20-APR-2006 23-FEB-2006. Synthetic. AEF94776;

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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked bhase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide drilo-cy3 which was used in the development of the novel process of the invention.
                                                                                Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; dna detection; DNA sequencing; DNA amplification; oligo dT40-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Optical DNA analysis process-related oligonucleotide dT40-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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                                                                                                                                             Example 5; Page 66; 94pp; German.
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  Hennig C;
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(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2006-185819/20.
                                          WPI; 2006-185819/20.
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modified_base
Cherkasov D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cherkasov D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AEF94782;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the sequence repeated as necessary. The solid phase is then washed and the sequence repeated as necessary. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process efficient and cheaper than prior art. Further claimed is present sequence is that of olisjonucleotide d740-biot which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                           Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; dna detection; DNA sequencing; DNA amplification; oligo dtT30-Cy3.
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1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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100.0%; Pred. No. 1.9
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                          Example 5; Page 66; 94pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= 5'-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEF94774 ståndard; DNA; 30 BP
               26-MAY-2004; 2004DE-10025695
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Matches 30; Conservative
                                                                                                                                      Cherkasov D, Hennig C;
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                                                                          (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
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(GENO-) GENOVOXX GMBH.
                                                     (CHER/) CHERKASOV D.
                                                                                                                                                                                   WPI; 2006-185819/20.
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modified_base
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Synthetic.
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Query Match

AEF94774;

RESULT 122 AEF94774/

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Gaps

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0; Indels

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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclid reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dT40-Cy3 which was used in the development of the novel process of the invention.
Example 2; Page 67; 94pp; German.
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DB 1; Length 30; 1.9e+02; Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other; .; 0 Query Match
1.1%; Score 30; DB
Best Local Similarity 100.0%; Pred. No. 1.9
Matches 30; Conservative 0; Mismatches

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Gaps

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Indels

2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738 30 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 1 쉽 ઠ

вв; dna detection; DNA sequencing; DNA amplification; oligo dtT30-Cy3. Optical DNA analysis process-related oligonucleotide dtT30-Cy3. AEF94758 standard; DNA; 30 BP 20-APR-2006 (first entry) AEF94758; RESULT 124 

Location/Qualifiers /\*tag= b /mod\_base= 5'-Cy3 Key modified\_base

Unidentified

Synthetic.

DE102004025694-A1

23-FEB-2006

26-MAY-2004; 2004DE-10025694

26-MAY-2004; 2004DE-10025694

(CHER/) CHERKASOV D. (HENN/) HENNIG C. (GENO-) GENOVOXX GMBH.

ΰ Hennig Cherkasov D,

WPI; 2006-185818/20.

This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a spiner-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex. German. Example 5; Page 67; 95pp;

co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solic phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present sequence is that of oligonucleotide dtT30-Cy3 which was used in the development of the novel process of the invention. 888888888888

Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;

ô Gaps . 0 Length 30; 0; Indels DB 1; Le 1.9e+02; 100.0%; Pred. .... Score 30; 1 Pred. No. 1.1%; Query Match Best Local Similarity 100. Matches 30; Conservative

2709 далалалалалалалалалалалалалалала 2738 ð 셤

Optical DNA analysis process-related oligonucleotide dT40-biotin. AEF94760 standard; DNA; 30 20-APR-2006 (first entry) AEF94760; AEF94760,

ss; dna detection; DNA sequencing; DNA amplification; oligo dT40-biot /mod\_base= 5'-Biotin-TEG Location/Qualifiers /\*tag= Key modified\_base Unidentified Synthetic

26-MAY-2004; 2004DE-10025694 DE102004025694-A1 23-FEB-2006. 

(GENO-) GENOVOXX GMBH. CHERKASOV D. HENNIG C. (CHER/) (HENN/)

26-MAY-2004; 2004DE-10025694

ΰ Cherkasov D, Hennig

WPI; 2006-185818/20.

Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex. Example 5; Page 67; 95pp; German.

This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present sequence is that of oligonucleotide dT40-biot which was used in the development of the novel process of the invention.

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30 AAAAAAAAAAAAAAAAAAAAAAAAA 1

(first entry)

AEF94766

Location/Qualifiers

/\*tag= b /mod\_base= 5'-Cy3

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Ultra-high parallel analysis process to analyse nucleic acid chains in
                                                                                                                                                                                                                              ss; dna detection; DNA sequencing; DNA amplification; oligo dtT30-Cy3.
                                                                                                                                                                                              Optical DNA analysis process-related oligonucleotide dtT30-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAY-2004; 2004DE-10025696.
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                                                                                         AEF94719 standard; DNA; 30
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(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2006-185820/20.
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                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                            20-APR-2006
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                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                            AEF94719;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                  ss; dna detection; DNA sequencing; DNA amplification; oligo dT40-Cy3.
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                                   DB 1; Length 30;
                                                                       Indels
Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                1.1%; Score 30; DB 1; Le
100.0%; Pred. No. 1.9e+02;
ative 0; Mismatches 0;
                                                                                                    2709 ААААААААААААААААААААААААААА
                                                                                                                        Location/Qualifiers
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/mod_base= 5'-Cy3
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                                                                                                                                                                                                                         AEF94766 standard; DNA; 30 BP.
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                                                                                                                                                                                                                                                                                             20-APR-2006 (first entry)
                              Query Match
Best Local Similarity 100.
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hennig C;
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GENOVOXX GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
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                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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Baeuml E;

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                                                                 This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide drillo-cy3 which was used in the development of the novel process of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 1.1%; Score 30; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 1.9e+02; Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
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Example 5; Page 95; 141pp; German.
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AEF94721/C
ID AEF947
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0; Gaps

Indels

1.1%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 1.9e+02;

100.0%; Pred. ...

Query Match
Best Local Similarity 100.0
Matches 30; Conservative

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Unidentified

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
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                                                                                 вв; dna detection; DNA sequencing; DNA amplification; oligo dT40-biot.
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                                                   Optical DNA analysis process-related oligonucleotide dT40-biotin.
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/mod_base= 5'-Biotin-TEG
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                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                          GENOVOXX GMBH
                                                                                                                                                                                                                                                                                                                                                              (CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2006-185820/20.
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Best Local Similarity
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                                                                                                                                                                 Key
modified_base
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                                                                                                                   Unidentified
                             20-APR-2006
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                                                                                                                                      Synthetic.
AEF94721;
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(GENO-)
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nuclectides are incorporated in the primer matrix by enzyme reaction, followed by are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nuclectides are detected and their co-ordinates logged and the signals removed. The marked nuclectides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dT40-Cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contryphan; leu-tryphan; anticonvulsant; neuroprotective; venom; cone snail; neurodegenerative disorder; epilepsy; neurotoxic injury; hypoxia; anoxia; ischaemia; stroke; cerebrovascular accident; brain trauma; spinal chord trauma; myocardial infarct; physical trauma; drowning; suffecation; perinatal asphyxia; hypoglycaemia; migraine; senile dementia; Alzheiner's disease; anyotrophic lateral selerosis; Parkinson's disease; Huntington's disease; bown's syndrome; PCR primer; korsakoff's disease; schizophrenia; neuronal damage; seizure; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                   Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conus stercusmuscarum contryphan PCR primer DHOG 496.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%; Score 30; DB 1; L. 1.00.0%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 95; 141pp; German.
                                                                                                                                                                                                                                                                                                                                                        Baeuml E;
                                                                                                                 /mod_base= 5'-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВЪ
                                                                                                                                                                                                                                                    26-MAY-2004; 2004DE-10025696
                                                                                                                                                                                                                    26-MAY-2004; 2004DE-10025696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX88521 standard; DNA; 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30; Conservative
                                                                                                                                                                                                                                                                                                                                                        Hennig C,
                                                                                                 *tag=
                                                                                                                                                                                                                                                                                                                       (GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                     (CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2006-185820/20.
                                                                                                                                                  DE102004025696-A1
                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                        Cherkasov D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-SEP-1999
                                                                                                                                                                                      23-FEB-2006
                                   Synthetic.
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The present invention relates to a method (M1) for detecting and/or isolating a nucleic acid having a homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) comprises treating a sample containing nucleic acid compounds with an locked nucleic acid (LNA) oligonucleotide (LO) to thereby detect and/or isolate a nucleic acid having the homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) is useful for detecting and isolating nucleic acids released from a lysed complex biological mixture comprising nucleic the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and/or isolating nucleic acid molecule having homopolymeric sequence or repetitive element or conserved nucleotide sequence involves treating sample containing nucleic acid compounds with locked nucleic
            /note= "10-mer deoxy-thymine and 5-mer non-base (t10-NB5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C-myc; cancer; HIV-1; AIDS; collagenase; Alzheimers disease; EGF; epidermal growth factor; GSTpi; HMGCOA; thallasemia; Herpes simplex virus; nerve growth factor receptor; globin; ss.
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                                                                                                                  AQ2, where AQ is anthraguinone"
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                                                                                                                                                                                                  /note= "Optionally LNA nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 35 BP; 0 A; 0 C; 0 G; 30 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence binding to and inhibiting the GSTpi gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2709 ААААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 23; Page 67; 104pp; English.
                                                                                                                                                           /*tag= c
/mod_base= OTHER
  /mod base= OTHER
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                                                                                                                                                                                                                                                                                                                     20-JUN-2003; 2003WO-IB006354.
                                                                                                                                                                                                                                                                                                                                                            24-JUN-2002; 2002US-0390928P.
                                                                              /*tag= a
/mod_base= (
/note= "5' ;
16. .35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           Jacobsen N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 85.7
les 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acid oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                 (EXIQ-) EXIQON AS
                                                                                                                                                                                                                                      WO2004020575-A2
                                                          modified base
                                                                                                                                      modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                           Kauppinen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
31-OCT-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ05003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence sequence represents a PCR primer for a contryphan peptides are lound in the venom of cone shalls. The contryphan peptides are useful as anticonvulsant agents, as neuroprotective agents, for managing pain, and for treating neuroprotective agents, for managing pain, and for treating neuroprotective agents, for managing pain, and for treating from an overstimulation of excitatory amino acid receptors. The contryphans are useful for the treatment and alleviation of excitatory amino acid receptors. The contryphans are useful for the treatment and alleviation of hypoxia, anoxia, or ischaemia which typicially follows strok, cerebrovascular accident, brain or spinal chord trauma, myocardial infarct, physical trauma, drownings, or spinal chord trauma, myocardial infarct, physical trauma, drownings, are further useful for the treatment of Alzheimer's disease, senile dementia, amyotrophic lateral sclerosis, Parkinson's disease, senile dementia, amyotrophic lateral sclerosis, Parkinson's disease, thutington's disease, bown's syndrome, Korsakoff's disease, senile associated with uncontrolled seizures. The contryphans are further useful for used prophylactically or to relieve the symptoms associated with a controlling pain and are effective in the treatment of migraine. They
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                                                                                                                                                                                                                                                                                              Grilley M;
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o. 2e+02;
0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection; isolation; locked nucleic acid; LNA; probe; ss.
                                                                                                                                                                                                                                                                                              Gray WR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 33 BP; 0 A; 1 C; 2 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                            Olivera BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2738
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100.0%; Pred. No. 2e+
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 20; 48pp; English.
                                                                                                                                                                                                                                                                                              Cruz LJ,
                                                                                                                                                                                                                                                                                                                                                                                             New pure contryphan peptides.
                                                                                                                                                     98WO-US026789
                                                                                                                                                                                          97US-0068737P
                                                                                                                                                                                                                                                    (UTAH ) UNIV UTAH RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30; Conservative
                                                                                                                                                                                                                                                                                            Jimenez E,
Hillyard DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .15
/*tag=
                                   Conus stercusmuscarum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LNA capture probe #3.
                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-419087/35.
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Best Local Simílarity
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modified_base
                                                                                                                                                     16-DEC-1998;
                                                                                                                                                                                          24-DEC-1997;
16-APR-1998;
                                                                        WO9933865-A1
                                                                                                                                                                                                                                                                                            Jacobsen R,
                                                                                                               08-JUL-1999
                                                                                                                                                                                                                                                                                                                  Watkins M,
              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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Gaps

Matches

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03-APR-2003; 2003US-00407818
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                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         misc RNA
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ADS19107/c
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                                                                                                                                                                                              Sequence forms triplex with the double stranded target sequence with G binding to G-C and T to A-T. The strand runs 3' to 5' in an antiparallel orientation and when targeted to a specific sequence will deactivate it. This allows for growth inhibition in cancerous cells, manipulation of cellular structural protein content; inhibition of IL-2 chain receptor; disbursing plaque formation in Alzheimer's disease; inhibiting EGF gene; modulating cholestrol synthesis through the HMGCAOA gene; suppressing NGF gene expression; arresting HSV-I replication and suppressing Beta-globin expression in thallassaemia and sicle cell anaemia patients. (Updated on 25-WAR-2003 to correct PA
                                                                                                                                              Synthetic oligo-nucleotide(s) which bind target duplex DNA - forming co-
linear triplex to control transcription process in gene-specific fashion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prion protein polymorphic microsatellite marker consensus sequence #25
                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; sheep;
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                                                                                                                                                                                                                                                                                                                                                DB 1; Length 29; 2.2e+02;
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s 0; Indels
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100.0%; Pred. No. 2.2
:ive 0; Mismatches
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                                                                                                                                                                              Claim 39; Page 30; 40pp; English
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                                                                        BAYU ) BAYLOR COLLEGE MEDICINE.
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                                                     88US-00287359.
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                               89EP-00313391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                       Hogan ME, Kessler DJ;
                                                                                                                           WPI; 1990-195509/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-215730/21.
                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
Matches 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microsatellite; ds
                                                                                    HOGA/) HOGAN M E.
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                                20-DEC-1989;
                                                     20-DEC-1988;
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           27-JUN-1990
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The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using a template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or compared pML. The method is used to identify microsatellite markers, in a consease and for predisposition to diseases associated with a predisposition to diseases and for predisposits of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, consensus or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a prion protein polymorphic microsatellite marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    labeling molecule, solubility, multisignal labeling reagent; 88;
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100.0%; Pred. No. 2.2e+02;
rative 0; Mismatches 0; Indels
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"Fluorescein labeled, 3' amine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 29 BP; 0 A; 0 C; 0 G; 29 T; 0 U; 0 Other;
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"Fluorescein labeled"
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"Fluorescein labeled"
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"Fluorescein labeled"
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"Fluorescein labeled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                        Page 50; 64pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADS19107 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-DEC-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29; Conservative
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/hote=
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/note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA-RNA hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-OCT-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a composition (I) of matter comprising an oligomer or polymer having two or more labeled groups, where the label or labels are chemically linked to the oligomer or polymer, one or more caractive groups, and one or more charged groups where the charged groups are covalently linked to the oligomer or polymer, or or on or more caractive groups, and one or more charged groups where the charged groups are covalently linked to the oligomer or polymer, or comprise part of the oligomer or polymer, or any of their combination. Also described are: a composition (II) comprising of their combination. Also been labeled using (I); and a composition (III) prepared by a target of labeling process comprising (I) providing a target for labeling, and a composition (II) providing a target for labeling, and a composition (II) providing a target for labeling, and a composition of (II) reacting the correct or (IV). (I) is useful for target and the labeling reagent to form the composition having the correcting or quantifying an analyte in a specimen. (II) or (III) is useful for (III), where the target is an analyte specific moiety, contacting the correcting or quantifying an analyte, which involves providing (II) or (III) where the target is an analyte specific moiety, contacting the amount of (II) or (III) bound to analytes in the specimen correspondent of (II) or (III) bound to analytes in the specimen or signal that is added to the analyte specific moiety, the containing the analyte with the presence of charged groups increases the presence of reactive groups analyte attachment of the multiple labeled groups increases the containing the analyte analyte analyte of the presence of charged group increases the containing the analyte analyte analyte and the presence of charged group increases the containing the analyte and the presence of charged group increases the containing the analyte analyte and the presence of charged group increases the containing the analyte analyte and the presence of charged group incre
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                                                                                                                                                                                                                                            Composition of multi signal labeling reagents, useful for detecting or quantifying analyte in specimen, has oligomer/polymer having labeled moleties, reactive groups and charged groups linked to oligomer/polymer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3'-amino oligonucleotide; solid support; benzene derivative; solid phase oligonucleotide synthesis; SPOS; controlled pore glass; CPG; bi-fluorescent probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3'-amino oligonucleotide #4 synthesised on a solid support.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%; Score 29; DB 1; Length 29; 00.0%; Pred. No. 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 29 BP; 0 A; 0 C; 0 G; 24 T; 5 U; 0 Other;
                                                                                                                                                           Donegan JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                             Example 6; SEQ ID NO 2; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                         Stavrianopoulos JG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADU07155 standard; DNA; 29 BP
                 03-APR-2003; 2003US-00407818.
                                                            RABBANI E.
STAVRIANOPOULOS J G.
DONEGAN J J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JAN-2005 (first entry)
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nes 29; Conservative
                                                                                                                                                                                                       WPI; 2004-727850/71.
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modified_base
                                                                                                                                                         Rabbani E,
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                                                                                       (STAV/)
(DONE/)
                                                               (RABB/)
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The invention relates to the preparation of 3'-amino oligonucleotide derivatives. The method comprises providing solid support benzene derivatives, synthesising an oligonucleotide pursuant to standard techniques for solid phase oligonucleotide synthesis (SPOS) where the oligonucleotide chain is assembled on the solid support, cleaving the oligonucleotide from the solid support and deprotecting the oligonucleotide completely except for the terminal protective group. The solid phase is a derivatised controlled pore glass (CPO). The 3'-amino oligonucleotides are useful in the synthesis of bi-fluorescent probes. This process suppresses the undesixed formation of unmodified 3'-OH oligonucleotides through cyclic phosphate intermediates and reduces the probability of errors resulting from the use of different reagents for different sets of oligonucleotides. This sequence represents and reduces the different sets of oligonucleotides. This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparation of 3'-amino oligonucleotide derivatives, useful to synthesis bl-fluorescent probes, comprises providing solid support compounds, synthesis an oligonucleotide chain is assembled on the solid support, cleavage and deprotection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; degenerative nervous system disease; graft versus host disease; Addison's disease; atrophic gastritis; desease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; multiple sclerosis; allergy; type II hypersensitivity, disease; type I hypersensitivity; disease; type II hypersensitivity; disease; type IV hypersensitivity; leprosy; infectious disease; viral infection; tingal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; breast cancer; des
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide synthesised in the examples of the present invention.
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Pred. No. 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Prec. ...
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/note= "Modified by NH2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 8; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADH70631 standard; DNA; 33 BP.
                                                                                                                                                                                                                                                        21-APR-2004; 2004US-00830484.
                                                                                                                                                                                                                                                                                                                                           21-APR-2003; 2003US-0464269P.
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                                                                                                                                                                                                                                                                                                                                                                                                                         (PROL-) PROLIGO LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-794500/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leuck M, Wolter A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                       JS2004220397-A1
                                                                                                                                                                     04-NOV-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH70631,
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17-OCT-2002

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The sequences given in ABQ80394-99 represent probes and targets which were used in the method of the invention for detecting a target betwee used in the method of the invention for detecting a target polynucleotide in a sample. The method comprises amplifying the target, polynucleotide in a sample. The method comprises amplifying the target, comparizing the target to oligonucleotide bound to nanoparticles in a nanoparticle detection system, determining the amount of signal generated as a result of binding, optionally repeating the above steps, and companies to the signal produced after at least one amplification cycle. The amount of signal produced after at least one amplification cycle. The companies the quantity of target polynucleotide in a sample, and for determining the quantity of target polynucleotide in a sample. The method is useful in research and analytical laboratories in DNA sequencing, in the field to detect the presence of specific pathogens, in the doctor's of fire for quick identification of an infection to assist in prescribing that are accounted in the method is based on observing colour change with the naked eye, hence the method is cheap, fast, simple, robust, do not require specialized or expensive equipment, and little or no
                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting target polynucleotide in a sample, by amplifying target, hybridizing it to oligonucleotides bound to nanoparticles in nanoparticle detection system, and determining amount of signal generated due to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gold nanoparticle conjugated APC gene probe SEQ ID NO 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 33 BP; 29 A; 1 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
1.0%; Score 27.2; DB 1;
Best Local Similarity 90.6%; Pred. No. 3.1e+02;
Matches 29; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2709 ААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  analyte detection; diagnosis; APC; probe; ss.
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                                                                                                                                                                                                                                                                                                                                Herrmann M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                 /*tag= a
/note= "Gold-S'-A"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 35; 74pp; English.
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                                                                                                                                                                              27-NOV-2002; 2002WO-US038069.
                                                                                                                                                                                                                                  30-NOV-2001; 2001US-0334644P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        instrumentation is required
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADX44838 standard; DNA; 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                Storhoff JJ, Fritz BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                 (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-617993/58
                                                                                        WO2003048769-A1
modified_base
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misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-APR-2005
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                                                                                                                                         12-JUN-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADX44838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADX44838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          요
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers or psecifically priming and allowing amplification of each 'Obeta gene, 'ObetaRNA or CDNA. The kit is useful for diagnosing organ transplant crafection and diagnosing and treating T-cell associated diseases including autoimmune diseases, degenerative nervous system diseases.

CC including autoimmune diseases, hypersensitivity diseases, infectious diseases, arrophic gastriitis. Degenerative nervous system diseases include multiple sclerosis and Alzheimer's disease. Hypersensitivity diseases include multiple clarosis and Alzheimer's disease. Hypersensitivities diseases include multiple clarosis and Alzheimer's disease. Hypersensitivities such as those present in CGoodpasture's syndrome and Type IV hypersensitivities such as those caused by viruses such as HVP, fungal infections such as those caused by the yeast ganus Candida, parasitic infections such as those caused by children in Neoplashic diseases include lymphoproliferative diseases include the harian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                      autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe, target, nanoparticle, detection; DNA sequencing; pathogen; infection; screening; colour change; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                Kit for diagnozing and treating T-cell associated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 28.8; DB 1; Length 33; ilarity 93.8%; Pred. No. 2.4e+02; Conservative 0; Mismatches 2. Tinals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 33 BP; 0 A; 0 C; 3 G; 30 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 АААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 825; 164pp; English.
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                                                                                                                                         99US-00263959
                                                                                                                                                                                           94US-00309335
95US-00531241
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                                                                                                                                                                                                                                                                                                                                       Rowen L;
                                                                                                                                                                                                                                                                (HOOD/) HOOD L E.
                                                                                                                                                                                                                                                                                      ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe APC 1-MUT.
                                                   US2002150891-A1.
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Local Si...
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    Homo sapiens
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ABQ80395;

# X S X & X E X B X S X X X B

Key

RESULT 137

ABQ80395

Query Match

Best Loca Matches

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Length 33; Indels

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Example 1; SEQ ID NO 2; 70pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conjugate probes.
                                                                                                                                                                                                                   Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-1986;
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26-MAY-1991
                                10-NOV-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUL-1987
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                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to the detection of a target analyte having at least two portions, which comprises forming a light scattering complex by contacting a sample containing specific binding complement with nanoparticle and with polysaccharide under conditions to allow binding of the specific complement to two or more portions of the target analyte and illuminating the light scattering complex under conditions to produce scattered light from the complex. The invention is useful for detecting target analyte, e.g. nucleic acids or proteins, useful for the diagnosis of genetic and infections diseases. The present sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "OTHER= Linked to gold-S' where S' indicates a
connecting unit prepared via an epiandrosterone disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analyte detection; DNA detection; protein detection; APC gene; probe; ss;
                                                                                                                                                                                                                                                                                                       Detection of target analyte, e.g. nucleic acids or proteins, useful for diagnosis of genetic and infectious diseases, comprises forming light scattering complex, and illuminating complex to produce scattered light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
/note= "5' conjugated to a gold nanoparticle via a
connecting unit"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 27.2; DB 1; Length 33; 90.6%; Pred. No. 3.1e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 33 BP; 29 A; 1 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human mutant APC 1 gene specific probe, APC 1-MUT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2740
                                                                                                                                                                                                                                               Bao YP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 AAAAAAAAAAAAAAAAAAAAAAAAAA 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APC probe conjugated to a gold nanoparticle.
                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 2; 70pp; English.
                                                                                                                                                                                                                                               Mueller UR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                 30-MAY-2003; 2003US-0474569P.
29-AUG-2003; 2003US-0499034P.
04-NOV-2003; 2003US-0517450P.
                                                                                                                      27-MAY-2004; 2004WO-US016656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AED67931 standard; DNA; 33
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/*tag≈ a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 90.67
Matches 29, Conservative
                                                                                                                                                                                                                                               Lucas A,
                                                                                                                                                                                                                (NANO-) NANOSPHERE INC
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                                                          WO2005008222-A2
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                                                                                                                                                                                                                                             Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                          from complex.
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Synthetic.
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                                                                                       27-JAN-2005
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Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light scattering complex formed.
                                                                                                                                                                                                                                                                                 Garimella V;
                                                                                                                                                                                                                                                                              Senical M,
                                                                                                                                                                                                                                                                                    Bao YP,
                                                                                                                                                                                                                                                                              Muller UR,
                                                  30-MAY-2003; 2003US-0474569P.
29-AUG-2003; 2003US-0499034P.
04-NOV-2003; 2003US-0517450P.
03-MAY-2004; 2004US-0567874P.
27-MAY-2004; 2004US-00854848.
22-NOV-2004; 2004US-00995051
                                                                                                                                                                                                                         (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                                                                 Lucas A,
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absence of a single target molecule or target analyte (e.g. nucleic acid, protein, libid, bacterium). The method involves contacting sample with one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is a human mutant APC gene specific probe. This sequence is used in the preparation of nanoparticle-oligonucleotide The present invention provides a method for detecting the presence or

1.0%; Score 27.2; DB 1; Length 33; 90.6%; Pred. No. 3.1e+02; Sequence 33 BP; 29 A; 1 C; 3 G; 0 T; 0 U; 0 Other;

Indels 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2740 0; Mismatches 29; Conservative

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Gaps

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Robertson JG;
                                                                                                                                                                                                                 Sequence of scissile link probe MRC071 (HL)
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                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                             86EP-00116906
                                                                                               AAN70281 standard; DNA; 27
                                                                                                                                                                                                                                                     Hybridisation; probe; ss
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7 10:41:34 2006

Tue Nov

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Query Match
               RESULT 141
                AAN70274/
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              g
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BP; 0 A; 0 C; 0 G; 21 T; 6 U; 0 Other; misc\_feature misc feature misc\_feature Sequence 27 25-MAR-2003 31-OCT-2002 25-APR-1990 WO8910415-A 02-NOV-1989. 27; Synthetic. AAN92240; 27 Query Match Duck P, Matches RESULT 14: AAN92240/ င္တ ò 셤 ö The patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.) NAZ ere noncomplementary nucleic acid sequences; ---S-- a scissilariamentary nucleic acid sequences; ---S-- a scissila linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolyaable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and Synthetic nucleic acid probes - comprising two nucleic acid sequences Synthetic nucleic acid probes - comprising two nucleic acid sequences linked by a scissile linkage. Gaps ö Length 27; 0; Indels Sequence 27 BP; 0 A; 0 C; 0 G; 25 T; 2 U; 0 Other; DB 1; Le 2.8e+02; Robertson JG; Sequence of scissile link probe MRC046 (PL) 0; Mismatches 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2735 Score 27; Pred. No. 27 AAAAAAAAAAAAAAAAAAAAAA 1 Crosby W, 1.0%; BP linked by a scissile linkage. 86EP-00116906 85US-00805279 Example; p29; 46pp; English. Example; p29; 46pp; English AAN70274 standard; DNA; 27 (revised)
(first entry) Conservative Hybridisation; probe; ss (MEIO-) MEIOGENICS INC. WPI; 1987-186567/27 Bender R, Best Local Similarity Matches 27; Conserv 04-DEC-1986; 05-DEC-1985; 03-OCT-2002 26-MAY-1991 08-JUL-1987 EP227976-A. Synthetic. AAN70274; Duck P,

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                                                       Gaps
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  Length 27;
                                                    Indels
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1.0%; Score 27; DB 1; Lv
100.0%; Pred. No. 2.8e+02;
iive 0; Mismatches 0;
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"deoxyribonucleotides."
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/note= "deoxyribonucleotides."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe MRCO46; solid support; ribonuclease.
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/*tag= b
/note= "ribonucleotides."
                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2735
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                                                                                                                                                                                                                                                                                          AAN92240 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                          Conservative
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/*tag=
/note= '
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Best Local Similarity
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                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SS probe MRCO46.
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ВР

27

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The sequence is given in the disclosure to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                        Replication of DNA - useful in genetic engineering and medical
                                                                                                                                     DNA sequence used in DNA replication method.
   27 AAAAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 20; 20pp; Japanese.
                                                                                                                                                                                                                                                            91JP-00240525.
                                                                                                                                                                                                                                                                                  91JP-00240525.
                                                                                                             (first entry)
                                                              AAQ40854 standard; DNA;
                                                                                                                                                                                                                                                                                                         (UYAR-) UNIV ARIZONA
                                                                                                                                                                                                                                                                                                                                 WPI; 1993-171830/21
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                                                                                                                                                                                                           JP05103673-A.
                                                                                                                                                                                                                                                         26-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                     applications
                                                                                                                                                                                                                                                                                  26-AUG-1991;
                                                                                                             23-SEP-1993
                                                                                                                                                                                                                                   27-APR-1993
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                                                                                                                                                                                     Synthetic.
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                                                                                      AAQ40854;
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    Gaps
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   Indels
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*tag= a
note= "deoxyribonucleotides."
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/note= "deoxyribonucleotides."
 Mismatches
                                                                                                                                                                                                                               Probe MRCO71; solid support; ribonuclease.
                        2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2735
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/note= "ribonucleotides.
18. .27
                                              27 AAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                        AAN92247 standard; DNA; 27
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 Conservative
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                                                                                                                                                         (revised)
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                                                                                                                                                                                                         3S probe MRC071,
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27;
                                                                                                                                                        25-MAR-2003
31-OCT-2002
                                                                                                                                                                                 25-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                      02-NOV-1989
                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                AAN92247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Duck P,
 Matches
                                                                                  RESULT 143
                                                                                                AAN92247
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                                                                                                                                                                                                                                                                                                    Vaccine, cytostatic; virucidal; bactericidal, fungicidal; anti-parasitic; immunostimulatory, tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                          Gaps
                                                          ;
0
                          1.0%; Score 27; DB 1; Length 27;
100.0%; Pred. No. 2.8e+02;
cive 0; Mismatches 0; Indels
Sequence 27 BP; 27 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                     2709 AAAAAAAAAAAAAAAAAAAAAAA 2735
                                                                                                     Immunostimulatory nucleic acid #822.
                                                                                                                                                                                      AAF99706 standard; DNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2000; 2000WO-US026383
                                                                                                                                                                                                                                              12-JUN-2001 (first entry)
                                                       27; Conservative
                                         Local Similarity
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Gaps

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2709 AAAAAAAAAAAAAAAAAAAAAA 2735

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                                                                                                                                                                                                                                                            response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immuno response. The present sequence is one such immunostimulatory nucleic acids can be pyrimidate an nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a ribt to a Thl immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anglogenesis inhibitor; ss; anglogenesis; solid tumour growth; tumour metastasis; grecancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial anglogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; anglofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                               Vaccinating against tumors, infectious diseases, allergies and asthmausing immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                              present invention relates to a method for stimulating an immune
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Angiogenesis inhibitory oligonucleotide #911.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ААААААААААААААААААААА 1
                                                                                                                                                                                                            Claim 101; Page 56; 338pp; English.
                                                                       Schetter C, Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (COLE-) COLEY PHARM GROUP INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-DEC-2001; 2001WO-US048458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-DEC-2000; 2000US-025534P
               (IOWA ) UNIV IOWA RES FOUND.
(COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABS78427 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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Best Local Similarity 100.0
Matches 27; Conservative
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                                                                                                             WPI; 2001-273485/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-DEC-2002
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                                                                           Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27
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                                                                                     The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Oster-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, hypertrophic scars. The present sequence is an antianglogenic nucleic
        Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbone; ss
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                                                                                                                                                                                                                                                                                                                                                                  Length 27;
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/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 27; DB 1; L
100.0%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunostimulatory nucleic acid SEQ ID NO: 842.
                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                             Claim 2; Page 35; 276pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL39406 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                     27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hartmann G;
                                                                                                                                                                                                                                                                                                      acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-154611/20
                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    developing cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200197843-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL39406;
                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 14
ABL39406/
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cancer, involving administering to actual marines by the venture developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal carcinoma, bladder cancer, brain and central nervous system (CNS) cancer, breakt cancer, concer, colon and rectum cancer, connective tissue cancer, ocsophageal cancer, colon and rectum cancer, connective cancer, leukaemia, live cancer, lung cancer, kidney cancer, larynx cancer, lawyloma, melanoma, myeloma, oral cavity cancer, ovarian cancer, pancreatic cancer, prostate cancer, rhaddomyosarcoma, skin cancer, stochach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the for treating or preventing exemplification of the invention

Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;

Gaps .; 0 Score 27; DB 1; Length 27; Pred. No. 2.8e+02; 0; Indels Score 27; 1.0%; Query Match Best Local Similarity 100.0 Warches 27; Conservative ò

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ABK66592 standard; DNA; 27 ABK66592; **ABK66592** 

02-JUL-2002 (first entry)

Human gene specific PCR primer #680.

Primer; 88; DNA microarray; differential expression analysis; human.

Homo sapiens

US6352829-B1

05-MAR-2002

99US-00225928 05-JAN-1999;

97US-00859998 21-MAY-1997;

(CLON-) CLONTECH LAB INC.

Bibilashvilli Jokhadze G, Chenchik A,

WPI; 2002-314699/35.

Producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.

Example 3; SEQ ID NO 680; 11pp; English.

The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of So distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where each gene specific primer has a sequence complementary to a distinct mRNA, and method is useful for producing a single gene specific primer. The method is useful for producing a sub-population of labeled NAs which is useful for analysing the differences in the RNA profiles between several different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the oppulations for each physiological source to identify differences in the population, where the comparison is preferably

; performed by hybridising the labeled NAs for each of the distinct physiological sources to an array of probe NAs stably associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO Gaps ö http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1 1.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0; Indels Sequence 27 BP; 4 A; 8 C; 6 G; 9 T; 0 U; 0 Other; Conservative 1 Similarity 27; Conserv Query Match Best Local S Best Loca Matches 

GAAGAGTCTACCCGACATAAGTCGAGG 1 B ACH03245 standard; DNA; 27 27 ACH03245; RESULT 149 ACH03245/c 셤 

2578 GAAGAGTCTACCCGACATAAGTCGAGG 2604

Immunostimulatory nucleic acid #880.

(first entry)

25-SEP-2003

Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.

Synthetic.

US2003050268-A1.

13-MAR-2003.

29-MAR-2002; 2002US-00112653.

29-MAR-2001; 2001US-0279642P.

(KRIE/) KRIEG A M. (BERG/) BERG D J.

Berg DJ; Krieg AM,

WPI; 2003-521815/49

Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.

English Disclosure; Page 32; 229pp; The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid

Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;

Query Match

1.0%; Score 27; DB 1; Length 27;

a "Allylamine modified uridine"

\*tag= 'note=

ocation/Qualifiers

c "Allylamine modified uridine"

note = "Allylamine modified uridine"

/\*tag=

/\*tag=

'note=

\*tag= d 'note= "Allylamine modified uridine"

e "Allylamine modified uridine"

/\*tag= /note=

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03-APR-2003; 2003US-00407818.
                                                                                                                                                                                                                                                  03-APR-2003; 2003US-00407818
                                                                                                                                                                                                                                                                     (RABB/) RABBANI E.
(STAV/) STAVRIANOPOULOS J G.
(DONE/) DONEGAN J J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                             US2004198971-A1
 DNA-RNA hybrid
                                                                                                                                                                                                               07-OCT-2004
                                                                                                                                                                                                                                                                                                         Rabbani E,
                   Synthetic
                                              misc_RNA
                                                                         misc_RNA
                                                                                                    misc RNA
                                                                                                                               misc_RNA
                                                                                                                                                          misc_RNA
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                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Multisignal labeling reagent associated oligonucleotide segid 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     labeling molecule; solubility; multisignal labeling reagent; ss;
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                                                                                                                                                                   ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 2.8e+02;
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            Indels
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  Pred. No. 2.8e+02;
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100.0%; Prec. ...
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                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 17; 221pp; English.
                                                                                                                                                  Immunostimulatory nucleic acid #822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВЪ
                                                                                            ADB37208 standard; DNA; 27 BP
                                                                                                                                                                                                                                                    02-FEB-2001; 2001US-00776479.
                                                                                                                                                                                                                                                                       03-FEB-2000; 2000US-0179991P.
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                                                                                                                                                                                                                                                                                                                             Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                 (first entry)
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ses 27; Conservative
            27; Conservative
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                                                                                                                                                                                                                                                                                                                                              WPI; 2003-657977/62.
                                                                                                                                                                                                                                                                                        (BRAT/) BRATZLER R
(PETE/) PETERSEN D
(FOUR/) FOURON Y.
    Best Local Similarity
                                                                                                                                                                                                                  US2003087848-A1.
                                                                                                                                                                                                                                                                                                                             Bratzler RL,
                                                                                                                                 04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-DEC-2004
                                                                                                                                                                                                                                   08-MAY-2003
                                                                                                                                                                                                Synthetic.
                                                                                                              ADB37208;
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Matches
                                                                           RESULT 150
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ADS19108/c
             Matches
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Donegan JJ;

Stavrianopoulos JG,

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The invention describes a composition (I) of matter comprising an oligomer or polymer having two or more labeled groups, where the label or coligomer or polymer, one or more cactive groups, and one or more charged groups where the charged groups reactive groups, and one or more charged groups where the charged groups are covalently linked to the oligomer or polymer or comprise part of the backbone of the oligomer or polymer, or any of their combination. Also described are: a composition (II) comprising a target molecule that has been labeled using (I); and a composition (III) prepared by a target of tabeling process comprising (I) providing a target for labeling, and a capture of the labeling reagent to form the composition having the target and the labeling reagent to form the composition having the community of the labeling an analyte in a specimen; and detecting or quantifying an analyte, which involves providing (II) or (III) where the target is an analyte, which involves providing (II) or (III) with a specimen suspected of containing the analyte and the analyte is an analyte (II) bound to analytes in the specimen contacting the analyte with the securing the analyte. (I) detects or quantifies analyte with this sensitivity. In (I), the multiple labeled groups increases the amount of signal that is added to the analyte specific moiety, the amount of signal that is added to the analyte specific moiety, the groups to a desirable target and the presence of charged group increases the solubility. This sequence represents a multisignal labeling reagent
                                                                                                        Composition of multi signal labeling reagents, useful for detecting or quantifying analyte in specimen, has oligomer/polymer having labeled moieties, reactive groups and charged groups linked to oligomer/polymer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 27 BP; 0 A; 0 C; 0 G; 22 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                       Example 7; SEQ ID NO 3; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 associate oligonucleotide.
WPI; 2004-727850/71.
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1.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 2.8e+02;

RESULT 152

ADU90227

Matches

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The present invention relates to a method for augmenting T-helper 1 cells
                                                                                                                                      Immunostimulant, Antiinflammatory, Antipsoriatic; Gastrointestinal-Gen., Antiulcer, Dermatological, Antiallergic, helper T-lymphocyte, immune stimulation, inflammation, psoriasis; inflammatory bowel disease, Crohns disease, ulcerative colitis; eczema; skin allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (i) is useful for treating non-allergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 27; DB 1; Length 27; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PR-1 promoter primer P41+ for in vivo footprinting.
                                                                                                                                                                                                                                                                                                                                                                              /note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;
                                                                                                    Immunostimulatory oligonucleotide, SEQ ID 880.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 880; 58pp; English.
                                                                                                                                                                                                                      contact dermatitis; 88; phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAR-2001; 2001US-0279642P.
29-MAR-2002; 2002US-00112653.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAY-2005; 2005US-00127654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV15487 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                    .. .27
*tag= a
                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (IOWA ) UNIV IOWA RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2005-768014/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  US2005250726-A1
                                                                                                                                                                                                                                                                                                Key
modified_base
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                                                                12-JAN-2006
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                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV15487;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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ID AAV1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose
      Gaps
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                                                                                                                                                                                                                                                                                                                          ss; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.
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      0; Indels
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                                                                                                                                                                                                                                                                                          Allergic response suppressor oligonucleotide #911.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;
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    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%; Score 27; DB
100.0%; Pred. No. 2.8
ive 0; Mismatches
                                        2709 AAAAAAAAAAAAAAAAAAAAAAAA 2735
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                                                                    AAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        an immunostimulatory nucleic acid.
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                                                                                                                                                                        ADU90227 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 153
AED75671/c
ID AED75671 standard; DNA; 27
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                                                                                                                                                                                                                                                    (first entry)
    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BRAT/) BRATZLER R L.
(PETE/) PETERSEN D M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-833006/82.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2004235774-A1.
                                                                                                                                                                                                                                                      10-FEB-2005
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27;
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                                                                                                                                                                                                            ADU90227
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Best Local &
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Synthetic.
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AAS00066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA-protein fusion; protein library; protein isolation; gene cloning; ss.
                                                                                                                                                                                                                                             Primer P41+ corresponds to nucleotides -735 to -706 relative to the transcription start site in the upstream region (see AAV15448) of the Arabidopsis PR-1 gene (see AAV15448). It was used in non-coding strand analysis of the PR-1 promoter region. In vivo footprinting analysis was performed of the PR-1 promoter region. Inducible in vivo footprints are located at positions -629 and -628 and at position -604 on the coding strand and at position -641 on the non-coding strand. The use of PR-1 promoter fragments to regulate gene expression in plants in the presence of chemical inducers is disclosed
                                                                                                                                                                                                   New chemically inducible promoter from Arabidopsis - used to regulate gene expression in response to e.g. salicylic acid.
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "attached to puromycin, a peptide acceptor"
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Promoter PR-1; salicylic acid, 2,6-dichloroisonicotinic acid; benzo(1,2,3)thiadiazole-7-carbothioic acid S-methyl ester; transgenic plant; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 27; DB 1; Length 29;
100.0%; Pred. No. 2.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                               Uknes SJ, Ward ER;
                                                                                                                                                                                                                                                                                                                                             Sequence 29 BP; 0 A; 2 C; 0 G; 27 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                     100.0%; Preu. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA-protein fusion oligonucleotide 30-P.
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                                                                                                                                                                                                                               Example 9; Page 32; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
                                                                                                                                                                Thorne L,
                                                                                                         97WO-US012626.
                                                                                                                          96US-0027228P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA94315 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
Warrhes 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                             (NOVS ) NOVARTIS CORP.
                                                                                                                                                               Lebel EG, Ryals JA,
                                                  Arabidopsis thaliana
                                                                                                                                                                                 WPI; 1998-120690/11.
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modified_base
                                                                    WO9803536-A1
                                                                                                         18-JUL-1997;
                                                                                                                           23-JUL-1996;
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                                                                                      29-JAN-1998
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The present sequence is one of a number of oligonucleotides which were used for the generation of RNA-protein fusions, including fusions having a myc epitope tag. The RNA-protein fusions comprise a protein covalently consisted to the 3' end of its own mRNA. This is accomplished by synthesis and in vitro or in situ translation of an mRNA molecule with a peptide acceptor attached to its 3' end. The RNA-protein fusions are incubated counder high salt conditions to produce a protein library. This method is cuseful for improving or altering existing proteins, as well as for isolating new proteins and nucleic acid or small molecule targets. It may can be used to improve human or humanised single-chain antibodies for the treatment of a number of diseases. The method is useful for the constitution of proteins with specific binding properties, for screening contractions. Unlike prior art, the new method does not rely on complex, which is very fragile and is therefore of limited use. The mathod does not rely on topological links between the protein and the conditions of the information of the protein is retained and can condition and can be recovered in readable, nucleic acid form
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                                                                                                                                                                                                                                         Producing protein or DNA libraries which are useful for improving existing proteins, by in vitro translating protein coding sequences to produce RNA-protein fusions and incubating these protein fusions under
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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/note= "Other= Covalently linked to puromycin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29 BP; 27 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%; Score 27; DB 1; L. 100.0%; Pred. No. 2.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure, Page 43; 121pp; English.
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99US-00247190.
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                                                            (GEHO ) GEN HOSPITAL CORP.
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                                                                                                                                                                                                                                                                                                                                            high salt conditions.
                                                                                                                            Roberts
                                                                                                                                                                                   WPI; 2000-533022/48.
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Matches 27; Conserv
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      09-FEB-1999;
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                                                                                                                         Szostak JW,
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AAK9863
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                                                                                                                                                                                                                                                      The sequence represents part of a branched encoding molecule used in methods to hybridise a capture probe to the addressing element of a DNA linker attached to am in vitro translated protein, in order to immobilise the protein to a solid support. The new methods are useful for tagging or encoding in vitro translated proteins with unique and minimal encoding concious and sorting these molecules onto solid supports. They are also useful for the identification of a desired binding partner. The method allows the use of pre-made sets of universal encoding molecules, such as nucleic acid(s) (analogues). These can be used in conjunction with nucleic acid(s) (analogues). These can be used in conjunction with corresponding universal microarrays or sets of microparticles to create new protein display systems which are flexible, modular, scalable and cost effective. The method allows the use of mucleic acid analogue which are not susceptible to enzymatic incorporation or polymerisation and are superior to conventional DNA/RNM. The proteins can also be labelled with fluorescent groups which can be used to monitor the protein in real time. The absence of RNA is advantageous as they can adopt secondary structures which are difficult to predict and can interfere with hybridisation steps and protein folding/function
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                                                                                                                                                                                      nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                      Encoding and sorting in vitro translated proteins, useful for the identification of desired binding partners, comprises attaching a racid linker to the protein and binding an encoding molecule to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detection; amplification; biomedical diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.0%; Score 27; DB 1; Length 29; 100.0%; Pred. No. 2.9e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 29 BP; 27 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAAAAAAAAAA 2735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C-myc epitope puromycin linker primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C-myc; epitope; detection; amplificat environmental monitoring; primer; ss.
                                                                                                                                                                                                                                   Example 3; Fig 9B; 48pp; English.
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                                               25-AUG-2000; 2000WO-US023414.
                                                                        99US-0151261P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100 les 27; Conservative
                                                                                                                                              WPI; 2001-183261/18.
 WO200116352-A1.
                                                                                              (PHYL-) PHYLOS
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                                                                        27-AUG-1999;
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                                                                                                                      Kuimelis RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                        08-MAR-2001
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                                                                                                                                                                                                            linker.
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Matches
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This invention describes a novel test system (A) which comprises at least one immobilized analyte (I) on an insoluble carrier and a polypeptide detection agent (II). specific for (I) and conjugated, via a linker, to an amplifier (III). (A) is used for direct, in vitro detection of (I) with amplification of the signal by polymerase chain reaction (PCR), or a related technique, applied to (III). The method is useful in biomedical diagnosis and environmental monitoring and can be used to detect a wide range of (I), e.g. diagnostic or pharmaceutical agents, secondary metabolites, herbicides or pesticides. (A) allow simultaneous, parallel detection of many different analytes (high throughput capacity).

The caltively simply (only a few incubation and washing steps are required) and with high sensitivity and selectivity. This sequence represents primer used in the ampification of the c-myc DNA fragment which encodes an epitope used to illustrate the method of the invention
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                                                                                                                                                                                                    System for detecting immobilized analyte, useful e.g. for biomedical diagnosis, has as detection agent specific polypeptide coupled to nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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1.0%; Score 27; DB 1; Length 29;
Best Local Similarity 100.0%; Pred. No. 2.9e+02;
Matches 27; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S cerevisiae alpha factor receptor STE2 vector linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 29 BP; 27 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
(AVET ) AVENTIS RES & TECHNOLOGIES GMBH & CO KG.
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/mod base= OTHER
                                                                                                                                                                                                                                           diagnosis, has as detection agacid for signal amplification.
                                                                                                                                                                                                                                                                                                                                        Example; Page 6; 12pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
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                                                                                                                                    WPI; 2001-381706/40.
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                                                                   Burgstaller P,
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modified_base
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Detecting and purifying biological material by (di)electrophoresis, useful e.g. for separating tissues and viruses, comprises using a probethat alters (di)electrophoretic properties.
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Example 1; Page 12; 20pp; German

The present invention relates to a method for the detection or purification of biological material by electrophoresis, which comprises (i) treating the biological material containing different species with a bioprobe and (ii) establishing an electric field for detection or purification of at least one complex formed between the biological material being tested and a specifically bound bioprobe. The method is wiruses, proteins, nucleic acids, lipids and/or other organic compounds. It can also be used for the isolation of specific bioprobes from a library of bioprobes. The present sequence is a linker described in the library of bioprobes. The present exemplification of the invention

Sequence 29 BP; 27 A; 2 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 1.0%; Score 27; DB 1; Length 29; 00.0%; Pred. No. 2.9e+02; 0; Indels 100.0%; Pred. No. 2.3 ive 0; Mismatches Best Local Similarity 100. Matches 27; Conservative Query Match

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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2735 

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AAV48087 standard; DNA; 30 BP. 27-OCT-1998 AAV48087; AAV48087 RESULT 

(first entry) Oligonucleotide 30-P. In situ translation; RNA-protein fusion; binding reagent; antibody; industrial catalyst; ss.

Synthetic.

Location/Qualifiers /\*tag= a /note= "Puromycin" Key modified\_base

WO9831700-A1

23-JUL-1998

97US-0035963P. 97US-0064491P. 14-JAN-1998; 21-JAN-1997; 16-NOV-1997;

98WO-US000807

(GEHO ) GEN HOSPITAL CORP.

RW, Liu Roberts Szostak JW,

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WPI; 1998-414032/35.

Selection of specific protein by screening protein-RNA fusions generated in vitro or in situ - useful for, e.g. identifying enzymes and antibodies with altered properties, potentially useful as catalysts or for therapy diaqnosis.

Disclosure; Page 39; 94pp; English.

The Oligonucleotides AAV48087, AAV48089-V48091 and AAV48096-V48098 and

the selection of a specific protein or RNA, by in vitro or in situ translations were used to generate RNA-protein or short or in situ translation of candidate RNA molecules to produce RNA-protein fusions, then selecting specific RNA protein fusions. The method is used to select proteins (or DNA encoding them) having altered properties, e.g. for identification of new binding reagents, to identify improved human antibodies or new enzymes. These proteins are potentially useful in diagnosis and therapy, or as industrial catalysts. The methods allow many counds of selection and amplification to be performed, resulting in enrichment of even very rare molecules and allowing isolation of proteins that bind specifically to almost any compound or catalyse almost any reaction 8.89999999999888

Sequence 30 BP; 27 A; 2 C; 0 G; 0 T; 0 U; 1 Other;

Gaрв ; 0 Length 30; 0; Indels Score 27; DB 1; Pred. No. 3e+02; 1.0%; Scc... 100.0%; Pred. No. cc... ... 0; Mismatches Best Local Similarity 100. Matches 27; Conservative Query Match

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RESULT 160 ADY75117

ADY75117 standard; DNA; 30

02-JUN-2005 (first entry) ADY75117;

Nucleic acid construct production associated polynucleotide #9.

recombinant DNA;

Unidentified

WO2005024018-A1

17-MAR-2005.

08-SEP-2004; 2004WO-JP013399.

08-SEP-2003; 2003JP-00315385

(ZOEG-) ZOEGENE CORP

Shiratori M; Sasaki T,

WPI; 2005-223378/23.

producing a nucleic acid construct, for producing a protein nucleic acid conjugate, comprises partial annealing a two single stranded nucleic acids, and coupling the nucleic acids to a third single stranded nucleic acid. 

Example 1; Page 30; 85pp; Japanese.

The invention describes a method of producing (M1) a nucleic acid construct having a first, second and third single stranded nucleic acid. The method comprises: coupling the first and third single stranded concerned by a chemical bond present on the same terminals of each nucleic acid by a chemical bond present on the same terminals of each cucleic acid; and coupling the second and third single stranded stranded nucleic acid by ligase. Also described are: a nucleic acid construct (I) nucleic acid by ligase. Also described are: a nucleic acid construct (I) obtainable by coupling the protein nucleic acid conjugate (II) obtainable by coupling the protein encoded by a coding sequence with (I); a DNA-RNA-coupling the protein encoded by a coding sequence with (I); and a double stranded DNA-protein conjugate obtainable by carrying (II); and a double stranded DNA-protein conjugate obtainable by carrying out a polymerase reaction with DNA and a degrading RNA of (III). (M1) is useful for producing a nucleic acid construct having a first, second and third single stranded nucleic acid. (I) Is useful for producing (II)

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which involves coupling protein encoded by (I) with (I) by a nucleic acid derivative. (I) Is translated with a protein translation system after heating (I) at 60 - 90 deg C and cooling (I) after the heating step. The method is useful for purifying (II) which involves separating and refining (II) by the labeling substance present on the third single of DNA or RNA that encodes a protein which involves selecting and purifying (II) and amplifying the coding sequence of (II). (II) Is useful for producing (III) which involves performing reverse transcription of mRNA of (II). (MI) Enables production of (I), which in turn enables efficient production of (II). This sequence represents a polynucleotide associated with the nucleic acid construct production method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mononucleotide repeat locus; human; BAT26; probe; microsatellite; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/mod_base= OTHER
/note= "Labelled with LightCycler fluorescent dye LC-Red-
640"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analyzing repeat sequences in DNA using a probe which hybridizes to adjacent repetitive and non-repetitive regions and determining hybrid melting point is useful to detect microsatellite instability such as in hereditary cancer.
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                                                                                                                                                                                                                                                                   Length 30;
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                                                                                                                                                                                                                                                                Score 27; DB 1;
Pred. No. 3e+02;
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                                                                                                                                                                                                                                                                1.0%;
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Best Local Similarity 96.4
Matches 27; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to detecting a mutant polynucleotide in a mixture of mutant polynucleotides, wild-type polynucleotides and unrelated polynucleotides, comprises using polymerase chain reaction amplification of extension products produced from mutant polynucleotide templates and by extension primers and probes. The method is useful in detecting mutated polynucleotides within a large population of wild-type polynucleotides. The method may be used in diagnosing or detecting cancer, such as colorectal cancer, in an individual. The present sequence
defined number of repeats, and determining the melting point temperature of the hybrid. The method is used to analyse microsatellites, especially microsatellite instability, particularly as a means for detecting hereditary tumours. Alternatively, the method is used to identify an individual in a population. The present sequence is a probe for Mononucleotide repeat locus BAT26, and was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mutant; wild-type polynucleotide; cancer; colorectal cancer; ss; probe.
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                                                                                                                                                                Length 32;
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                                                                                                                             Sequence 32 BP; 27 A; 1 C; 2 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                               Score 27; DB 1; Le Pred. No. 3.1e+02;
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Pred. No. 3.1e+02;
                                                                                                                                                     1.0%; Scc...
100.0%; Pred. No. ...
... 0; Mismatches
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                                                                                                                                                                                            27; Conservative
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                                                                                                                                                                               Local Similarity
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                                                                                                 invention
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                                                                                                                                                                                            Matches
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The present invention relates to an immunostimulatory composition, which comprises at least one oligonucleotide comprising both an RNA region and a DNA region. The composition is useful for enhancing an immune response or inducing cytokines. It can be useful for enhancing an immune response treating diseases, including pathogenic infection, (non-)malignant tumnours (e.g. cancers of the brain, lung, ovary, breast, prostate or colon, or carcinomas and sarcomas), autoimmune diseases or allergic finitis, hay fever or food allergies), Lyme disease, hepatitis, HIV or malaria. The composition is also useful for treating, preventing or ameliorating the symptoms resulting from exposure to a biowarfare agent, e.g. Ebola, Anthrax or Listeria. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New immunostimulatory compositions comprising RNA/DNA hybrid oligonuclectides, useful for enhancing an immune response or inducing cytokines, particularly for treating diseases, e.g. cancer, allergy or HIV infection.
                                                                                                                      Phosphorothioate substituted RNA/DNA hybrid oligonucleotide SEQ ID NO:
                                                                                                                                                                     infection; allergy; cancer; hypersensitivity; bio-warfare; immunostimulant; antiallergic; cytostatic; antimicrobial; anti-HIV; immunosuppressive; protozoacide; virucide; hepatotropic; gene therapy; antiinflammatory; antibacterial; ss
                                                                                                                                                      hybrid; phosphorothioate backbone; immunostimulatory; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 28 A; 1 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     /note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 30; 68pp; English.
                                                                                                                                                                                                                                                                                                                       /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Klinman DM;
                              BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-2001; 2001WO-US018276.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-2000; 2000US-0209797P
                              ABL35101 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                   Ω
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                      1. .14
/*tag= b
17. .30
/*tag= c
                                                                                                                                                                                                                                                                                                        1. .14
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BIOS-) BIOSYNEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-130570/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Flora M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200193902-A2
                                                                                                                                                                                                                                                                                          Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-2001
                                                                                               04-APR-2002
                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mond JJ,
                                                               ABL35101;
                                                                                                                                                                                                                                                                                                                                                                       misc_RNA
                                                                                                                                                                                                                                                                                                                                                                                                       misc_RNA
RESULT 163
                ABL35101
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This invention describes a novel method for the selective detection of nucleic acids which comprises amplification of the nucleic acid with the help of labeled primers and detection with a trap probe. The methods and reagents are used for the detection of a marker primer and at least 2 immobilized (or immobilizable) trap probes with the corresponding nucleic acid sequence of interest for mutation analysis. The method can be used to detect a specific sequence in a sample of one or more nucleic acids by using several sets of primers and trap probes (i.e. in an array). The methods are useful in molecular biology and disgnostic applications, especially for simultaneous detection of multi-pathogens, typing of organisms, analyzing genetic diversity and sequencing of genes or genomes. This sequence represents a primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Coxsackie virus strain VD2921; diabetogenic coxsackie B virus-4; CBV-4; strain VD2921; VP1; VP2; VP3; VP4; P2A; P2B; P2C; P3A; P3B; P3C;
                                                                                                                                                    RNA polymerase; rpo-beta; detection; diagnostic; trap probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                          Selective detection of nucleic acids by amplification with labeled primers and detection with a trap probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Coxsackie B virus 4 (CBV-4) strain VD2921, PCR primer dT26V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 26.6; DB 1; Length 27; 96.3%; Pred. No. 3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 27 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    M. tuberculosis rpo-beta primer 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1c; Page 19; 27pp; German.
                                                                                                                                                                                                                                                                                                                                                  (HOFF ) ROCHE DIAGNOSTICS GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                      99EP-00110458.
                                                                                                                                                                                                                                                                                                                    98DE-01024900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                        Mycobacterium tuberculosis
                             AAZ43904 standard; DNA; 27
                                                                                             10-MAR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                   Brand J;
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-055287/05.
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Best Local Similarity
Matches 26; Conserv
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                                                                                                                                                                                                                                                                                      29-MAY-1999;
                                                                                                                                                                                                                                                                                                                       04-JUN-1998;
                                                                                                                                                                                                                        EP962536-A1
                                                                                                                                                                                                                                                        08-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                   Weindel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABX12469;
                                                               AAZ43904;
RESULT 164
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                AAZ43904
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Gaps ö

Score 26.8; DB 1; Length 30; Pred. No. 3.18+02; 0; Mismatches 2; Indels

Query Match 1.0%; Best Local Similarity 93.3%; Matches 28; Conservative 0

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The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: method of determining (M2) microsatellite markers (MM) for a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or concerned pML, and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a diseases and for prediagnosis of such diseases especially prion diseases. Cut also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, companient of particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion companient (PTP) comprising a polymorphic microsatellite locus.
                                                                                                                                        Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic nucleic acid probes - comprising two nucleic acid sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26.2; DB 1; Length 31; Pred. No. 3.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 31 BP; 0 A; 3 C; 0 G; 28 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence of scissile link probe MRC060 (HL)
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                                                                                                                                                                                                                    Example 3; Page 28; 64pp; German.
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                                                                           Han
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Crosby W,
     09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86EP-00116906
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28; Conservative
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                                                                          Geldermann H, Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MEIO-) MEIOGENICS INC.
                                        (UYHO-) UNIV HOHENHEIM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
                                                                                                             WPI; 2004-215730/21
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1987-186567/27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-OCT-2002
26-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUL-1987.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                          primers for detecting nucleic acids from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
diabetes; diabetogenic enterovirus; beta cell loss; blindness; renal failure; leg amputation; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.0%; Score 26.2; DB 1; Length 27; 96.3%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                           the diabetogenic coxsackie B virus-4 strain VD2921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 27 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cow prion protein microsatellite locus primer #82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 44; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     microsatellite; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                         Polymerase chain reaction and
                                                                                                                                                                                                                               (INNO-) INNOVENTUS PROJECT AB
                                                                                                                                                             19-JUN-2002; 2002WO-IB003278
                                                                                                                                                                                              20-JUN-2001; 2001SE-00002198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2002; 2002DE-01036711.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 96.3
Matches 26; Conservative
                                                                                                                                                                                                                                                                    Frisk GE,
                                                                                                                                                                                                                                                                                                      WPI; 2003-278229/27.
                                                                                      WO2002103060-A2
                                                   Coxsackievirus.
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Gaps

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3; Indels

a scissile linkage.

linked by

Length 26;

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The patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- = a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule in the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.)
                                                            The patent claims a new molecule of formula (NA1----S----NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S-- = a scissile linkage. n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be FL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support) or ML (Hydrolysable Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic nucleic acid probes - comprising two nucleic acid sequences linked by a scissile linkage.
                                                                                                                                                                                                                                                                                                DB 1; Length 26; 3.2e+02;
                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                               Sequence 26 BP; 0 A; 0 C; 0 G; 22 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robertson JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         scissile link probe MRC059 (HL)
                                                                                                                                                                                                                                                                                                Query Match 0.9%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 3.2 Matches 26; Conservative 0; Mismatches
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                                  Example; p29; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAN70275 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1987-186567/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-1986;
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26-MAY-1991
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                                                                                                                                                                                                                                field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 168
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Sequence 26 BP; 0 A; 0 C; 0 G; 22 T; 4 U; 0 Other;

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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                            Gaps
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0.9%; Score 26; DB 1; Length 26;
Best Local Similarity 100.0%; Pred. No. 3.2e+02;
Matches 26; Conservative 0; Mismatches 0; Indels
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0.9%; Score 26; DB 1; Le
100.0%; Pred. No. 3.2e+02;
iive 0; Mismatches 0;
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/*tag= b
/note= "ribonucleotides."
15. .26
/*tag= c
/note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                     "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                    Probe MRCO59; solid support; ribonuclease.
                                                      2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2734
                                                                                 Location/Qualifiers
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                                                                                                                                                                                                               (revised)
(revised)
(first entry)
 Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                          AAN92241 standard; DNA;
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bender R;
                                                                                                                                                                                                                                                                         SS probe MRCO59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-APR-1988;
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                                                                                                                                                                                                                                                                                                                                                                          misc_feature
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25-APR-1990
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                                                                                                                                                                                                                                                                                                                                                              Key
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BP.

(first entry)

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cDNA library production; SCLA; gene chip technology; differential screening; pathological diagnosis; genetic identification; single-cell cDNA library amplification; ds.
                                                                                                                                       CDNA library production method related oligonucleotide SEQ ID NO:
                                                 AAF77536 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-243448/25.
                                                                                                                                                                                                                                                                                                                            20-NOV-1998;
                                                                                                                                                                                                                                                                 US6197554-B1
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                                                                                                           23-MAY-2001
                                                                                                                                                                                                                                    Synthetic.
                                                                            AAF77536;
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(YING/)
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                   RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3' end. It is used by reacting excess probe with a target mucleic acid; form 2 or more probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. The probe system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Probe MRCO60 is bound by a hydrolysable linkage to a solid support at its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 26 BP; 0 A; 0 C; 0 G; 22 T; 4 U; 0 Other;
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"deoxyribonucleotides."
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/note= "deoxyribonucleotides."
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Best Local Similarity 100.0%; Fred. No. 3.2.

Best Local Similarity 0; Mismatches
                                                                                                                                                                                                                                                            Probe MRCO60; solid support; ribonuclease.
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/*tag= b
/note= "ribonucleotides."
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               26 AAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                       AAN92242 standard; DNA; 26
                                                                                                                                                                                                 (first entry)
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/*tag= c
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*tag=
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                                                                                                                                                                  (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bender R;
                                                                                                                                                                                                                               SS probe MRCO60.
                                                                                                                                                                                                                                                                                                                                     misc_feature
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31-OCT-2002
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                                                                                                                                    AAN92242;
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for use

98US-00197951. 98US-00197951.

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Ying

Chuong C,

LIN S. CHUONG C. YING S.

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The present invention describes a method of producing full-length cDNA libraries from single cells, designated single-cell cDNA library amplification (SCLA). The method is useful in gene chip technology, differential screening, pathological diagnosis, physiological prognosis and genetic identification. No further information about this sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; phosphodiesterase; PDE; zcytor13; antiasthmatic; antiarthritic; antipsoriatic; cytostatic; antiatherosclerotic; antinfertility; cardiant; antinflammatory; dermatological; wound healing; antiviral; antibacterial; therapy; inflammatory bowel disease; diverticulitis; spermatogenesis; sperm capacitation; immunocontraceptive; vaccine; cancer; reperfusion ischaemia; psoriasis; melanoma; myocarditis; PID;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
Generating a complete full-length cDNA library from single cells for in gene chip technology, involves reverse transcribing intracellular mRNAs, adding polynucleotide tail and amplifying formed cDNAs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human full length zcytor13 cDNA isolating polyA PCR primer, ZC7764b.
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                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                    Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
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                                                                                                              Disclosure; Col 11-12; 11pp; English.
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Gaps

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0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02;

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The patent discloses novel human phosphodiesterase (PDE), zcytor13 cDNA and its corresponding protein. Zcytor13 protein is used to promote wound tealing in tissues, to exhibit anti-bacterial and anti-viral effects and to identify modulators (e.g. agonists or antagonists). Zcytor13, its action as are useful in the treatment of inflammatory heart or cardiovascular conditions, muscle inflammation, inflammation, heart or cardiovascular conditions, muscle inflammation, inflammation, as and after surgary, arthritis asthma, inflammation, inflammation, as immunocontraceptive or anti-fertility vaccine and for treating male infertility. Zcytor13 protein and its antibodies are used to diagnose cancer, reperfusion ischaemia, asthma, psoriasis and melanoma. Zcytor13 corteins are used to enhance fertilisation. Zcytor13 antagonists are used to cracer, reperfusion ischaemia, pelvic inflammatory disease. [PID), corteins are used to enhance fertilisation. Zcytor13 antagonists are used to ensure and other inflammatory disease. [PID), corteins are used to enhance fertilisation. Zcytor13 antagonists are used to sasociated with vasoconstriction, heart arrhythmia, congestive heart disease, muscle spasms and fatigue. They are used for detecting human chromosomal abnormalities. Zcytor13 cDNAs are used in gene therapy.

C Zcytor13-cytokine fusion proteins or antibody-cytokine fusion proteins care useful for enhancing in vivo killing of target tissue. The present csquence is a polyA PCR primer, Zc7764b which is used to isolate full
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                  Novel human phosphodiesterase polypeptide, zcytor13 and polynucleotide encoding it, for detecting human chromosomal abnormalities, identifying modulators and treating inflammatory and cardiovascular diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
pelvic inflammatory disease; eczema; scleroderma; vasoconstriction;
heart arrhythmia; congestive heart disease; muscle spasm; fatigue;
chromosomal abnormality; gene therapy; PCR primer; ss.
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Best Local Similarity luv...
Local 26; Conservative
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                                                                                                                                                                                                                                                                                  (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                            Presnell SR, Novak JE,
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-266312/27
                                                                                                                         WO200125444-A2
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                                                                                    Homo sapiens
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Human; zsig63; chromosome 4q12-4q13; salivary protein; antimicrobial; ss; microbial infection; tooth decay; periodontal disease; thrush; emphysema; gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; epithelial wound; chronic tissue damage; cystic fibrosis; acquired immunodeficiency syndrome; AIDS; lung infection; sarcoidosis;
                                                                                                                                                                                                                                            Generating amplified messenger RNA sequences from single cells, involves cycling steps of reverse transcription, denaturation, double-stranded DNA sequences and in vitro transcription.
                                                                                                                                                                                                                                                                                                                                  RNAs with
                                                                                                                                                                                                                                                                                                                                 The present invention relates to generating amplified messenger RNAs with polymerase reaction activity, comprising cycling steps of reverse transcription, denaturation, double-stranded cDNA synthesis and in vitro transcription. The invention is used for generating amplified mRNAs from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chronic bronchitis; gene therapy; protein therapy; primer; 2C7764a
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 0.9%; Score 26; DB 1; Length 26; Local Similarity 100.0%; Pred. No. 3.2e+02; les 26; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human zsig63 cDNA sequencing primer ZC7764a.
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                                                                                                                                                                                            Chuong C, Widelitz RB;
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                                                                                                                                                                                                                                                                                                        Disclosure; Page 17; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                           limited mRNAs from single cells
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                                                                 99WO-US012461
                                                                                              99WO-US012461
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                                                                                                                                       (YING/) YING S.
(CHUO/) CHUONG C.
(WIDE/) WIDELITZ R B.
                                                                                                                                                                                                                      WPI; 2001-061734/07.
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                                                                                                                                                                                              Ying S,
           WO200075356-A1
                                                                                                                          LIN S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                  04-JUN-1999;
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Tue Nov

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Polynucleotides encoding salivary proteins useful as anti-microbial
                                           agents.
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Example 1; Col 53; 29pp; English

region of human chromosome 4 and encoding a zsig63 polypeptide, a microbial activity. Due to their microbial activity. Due to their microbial activity, the sequences can be used in the study of microbial infections, e.g. for recombinant production of anti-microbial proteins. The sequences can be used in the treatment of tooth decay, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, spitchellal wounds, chronic tissue damage, acquired immunodeficiency syndrome (ADDS), cystic fibrosis, lung infections, sarcoidosis, emphysema and chronic bronchitis. This sequence represents a sequencing primer for CDNA encoding human zsig63 The invention relates to a polynucleotide derived from the 4q12-4q13

Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;

0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; 0; Indels 100.0%; Prec. ... 2708 TAAAAAAAAAAAAAAAAAAAAAAAA 2733 26; Conservative Local Similarity Query Match Matches ò

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Gaps

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26 TAAAAAAAAAAAAAAAAAAAA 1 ABS52638 standard; DNA; 26 BP. (first entry) 15-NOV-2002 ABS52638; RESULT 175 ABS52638/c 셤

Human; secreted salivary protein; zsig63; immunogen; zsig63-cytokine; antibody-cytokine; in vivo killing; pathological microbe; bacteria; fungal; viral; infection; salivary gland; anti-microbial; dental caries; tooth decay; periodontal disease; thrush; gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; microflora; epithelial wound; pathogenic colonisation; invasion; pro-inflammatory; ehrompetent immune system; diabetes; anti-inflammatory; incompetent immune system; AIDS; acquired immunodeficiency syndrome; chemocherapy; radiation treatment; lung infection; cystic fibrosis; digestion; PCR; primer; ss.

Human secreted salivary protein zsig63 PCR primer ZC7764a.

Homo sapiens.

JS2002081701-A1.

27-JUN-2002.

03-AUG-2001; 2001US-00922480.

17-MAR-1999; 99US-0124820P.

(ADLE/) ADLER D A. (SHEP/) SHEPPARD P O.

Adler DA, Sheppard PO;

WPI; 2002-635468/68.

Novel secreted salivary protein, zsig63 and polynucleotide encoding it useful for treating microbial infections, inflammatory conditions, dental caries and lung infections associated with cystic fibrosis.

Example 1; Page 29; 33pp; English

The present invention relates to a new secreted salivary protein, zsig63. The invention is useful for detecting in a test sample, the presence of an antagonist or agonist of zsig63 protein activity. The invention is also useful as an immunogen for producing an antibody to zsig63 polypeptide. zsig63-cytokine fusion proteins or antibody cytokine fusion proteins or antibody-cytokine fusion proteins or antibody-cytokine fusion proteins or antibody-cytokine fusion protein are useful for enhancing in vivo killing of target tissues. The manaceutical composition comprising purified zsig63 polypeptide are useful in the treatment of conditions associated with pathological microbes, including bacterial, fungal and viral infections. High polypeptides are useful for treatment of dental caries (tooth decay), periodontal disease, thrush and gastrointestinal disease. Other prevention of infection in winnary tract infections, vaginal infections, prevention of infection in skin and other epithalial wounds. The polypeptides can be used to establish normal microflora and protect against pathogenic colonisation and invasion. The invention is useful when pro-inflammatory activity is desired. Applications for such pro-inflammatory activity is desired. Applications for such pro-inflammatory activity is desired. Applications for such pro-inflammatory activity is almitted or damaged vascular system e.g., damage in extremities associated with diabetes. Antagonists to zsig63 cuseful for the treatment of pathents having incompetent immune system, and purch as and contribations. such as AIDS (acquired immunodeficiency syndrome) patients amuno system; that have undergone chemotherapy, radiation treatment. The invention is also useful for the treatment of lung infections associated with cystic fibrosis and its agonists or antagonists are useful for aiding digestion. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention for identification of zsig63 

Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;

Gaps ; 0 0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; tive 0; Mismatches 0; Indels Query Match 0.99 Best Local Similarity 100. Matches 26, Conservative

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ABK66591 standard; DNA; 26 ABK66591; **ABK66591** 

RESULT 176

ВP

(first entry) 02-JUL-2002 Human gene specific PCR primer #679.

Primer; ss; DNA microarray; differential expression analysis; human.

Homo sapiens

US6352829-B1

05-MAR-2002.

99US-00225928. 05-JAN-1999;

97US-00859998. 21-MAY-1997;

(CLON-) CLONTECH LAB INC

Bibilashvilli R; Jokhadze G, Chenchik A,

WPI; 2002-314699/35.

Producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.

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                      The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological acids (NAs) comprising contacting a NA sample from a physiological conditions to enzymatically generate sub-population of NAs, where each conditions to enzymatically generate sub-population of NAs, where each labeled nAs is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled NAs which is cuseful for analysing the differences in the RNA profiles between several different physiological sources, where the method comprises producing cuspulation of labeled NAs for the different physiological sources, comprising the populations for each physiological source to identify differences in the population, where the comparison is preferably comprising the population, where the comparison is preferably difference by physiological sources to an array of probe NAs stably associated with the surface of a substrate to produce a hybridisation pattern for each of the sources, where differential gene expression assays are utilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal cypecasion analysis of diseased a normal tissue e.g. neoplastic a normal cypecasion analysis of diseased a normal tissue e.g. neoplastic and the sequence data for this patent did not form part of the printed the sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from USPTO at http.wipo.seqdata.uspto.gov/sequence.html?pocID=635282981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; secreted salivary protein; zsig63 protein; host defense protein; immune modulating factor; antipathogenic; cell-cell signalling molecule; growth factor; cytokine; growth factor hormone activity; dental carries; infection; tooth decay; periodontal disease; gastrointestinal disease; thrush; urinary tract infection; vaginal infection; diabetes; obesity; anti-inflammatory; chronic tissue damage; lung dysfunction; restenosis; gene therapy; salivary gland dysfunction; prostate gland dysfunction; forensic DNA profiling; chondrosarcoma; atherosclerosis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZC7764a primer used in the identification of human zsig63 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 26 BP; 7 A; 6 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2360 AGCAAGGGTACGCTGGGCAAGTTCAC 2385
   Example 3; SEQ ID NO 679; 11pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 AGCAAGGGTACGCTGGGCAAGTTCAC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD45055 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAR-1999; 99US-0124820P.
17-MAR-2000; 2000US-00527345.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-AUG-2001; 2001US-00923236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-DEC-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adler DA, Sheppard PO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ADLE/) ADLER D A. (SHEP/) SHEPPARD P O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-642378/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2002090677-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD45055;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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The invention relates to human secreted salivary polypeptide designated as zaig63 and nucleic acid molecules encoding such polypeptides. Zaig63 can be used in detecting agonists and antagonists of its acityity, and is also useful as a host defense polypeptide. immune modulating factor, antipathogenic polypeptide, call-cell signalling molecule, growth factor, cytckine, or as secreted extracellular matrix associated protectins with growth factor hormone activity. It is useful for treating conditions associated with pathological microbes, including bacterial, fungal and viral infections, for treating dental carries (tooth decay), periodontal disease, thrush and gastrointestinal disease, for treating urinary tract infection, vaginal infection and for preventing infection in skin and cother epithelial wounds. zsig63 is useful for establishing normal cother epithelial wounds. zsig63 is useful for establishing normal cother epithelial wounds. zsig63 is useful for establishing normal cother pathological minipage e.g. damage in extremities associated with disease and useful as anti-inflammatory agents. It is useful as a marker of lung dysfunction, salivary gland dysfunction, or dysfunction of prostate gland. It is also therapeutically useful for aiding digestion. Polymuclectides of the invention are used in gene therapy for increasing cor inhibiting zsig63 activity, for detecting abnormalities on human chromosome 4 associated with disease or other human traits and as consecuted in forensis DNA profilling. Sequences of the invention are useful for stimulating proliferation or differentiation of adipocytes and for monormal sequences. It is a second to a second or differentiation of and or and proved the province as a second or servince or inhibiting chondrosarcoms, atherocleories, restenoslas and observe. The monormal sequences of the province of a second or experience or the invention are second or experiences and provinces of the invention or experiences. The second or experiences of the provinces of the province or an 
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Novel secreted salivary polypeptide, zsig63, useful as antimicrobial agent for treating microbial infection, dental carries, periodontal disease, thrush gastrointestinal disease, and for aiding digestion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; B-cell proliferation; anti-tumour response; immune system; immunostimulant; cytostatic; human; sequencing primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
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100.0%; Pred. No. 3.2e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human zalphall Ligand sequencing primer ZC7764a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2708 TAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                                                                                       Example 1; Page 30; 33pp; English
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Best Local Similarity 100.C
Warshes 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6307024-B1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-OCT-2001.
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99US-0123547P. 99US-0123904P. 99US-0142013P.

> 11-MAR-1999; 01-JUL-1999;

09-MAR-1999;

(ZYMO ) ZYMOGENETICS INC.

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Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                             2708 TAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                    TAAAAAAAAAAAAAAAAAAA 1
                                                                 Example 7; Col 139; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                            (GENO-) GENOMIC EXPRESSION APS
                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                            100.08;
                                                                                                                                                                                                                                                                                                                                                                                        24-JAN-2001; 2001DK-00000126.
12-FEB-2001; 2001US-0267704P.
                                                                                                                                                                                                                                                                                                                                                                              24-JAN-2002; 2002WO-DK000052
                                                                                                                                                                                                                                                                       AAD43853 standard; DNA; 26
                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                  Conservative
      Presnell SR,
Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-636542/68.
                        WPI; 2002-040208/05
                                                      antigenic response
                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                     WO200259357-A2.
                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                 26;
                                                                                                                                                                                                                                                                                                                                                                                                                        Pedersen ML;
                                                                                                                                                                                                                                                                                               14-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                01-AUG-2002.
                                                                                                                                                                                                                                                                                                                              primer; ss
      Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                  AAD43853;
                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                       Single
                                                                                                                                                                                                            Best Loca
Matches
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polynucleotide tag from a biological sample by cleaving one of the complementary strands of a double stranded polynucleotide with a cleavage agent capable of recognising a double stranded polynucleotide comprising complementary strands and cleaving only one of the strands of the polynucleotide in the process of generating a single stranded polynucleotide tag. The method is useful for separating, analysing, quantifying or obtaining single stranded polynucleotides comprising originating partly, and preferably wholly from a source of DNA and/or RNA in a sample comprising biological cells. The method is particularly for analysing gene expression (expression profilling or differential gene expression), or in diagnosing clinical conditions. The present sequence is a primer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunostimulant; oligodeoxynucleic acid; ODN; vaccine; DNA-RNA hybrid;
for analyzing gene expression or diagnosing clinical conditions, comprises employing nicking endonucleases that cleave complementary
                                                                                                                          The invention relates to a method for obtaining a single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 26; DB 1; Lo
Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "thiophosphate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INTE-) INTERCELL BIOMEDIZINISCHE FORSCHUNGS. (CIST-) CISTEM BIOTECHNOLOGIES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. wv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAAAAAAAAAAAA 2734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligodeoxynucleic acid molecule ODN 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26 AAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                  Example; Page 294; 302pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schmidt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-MAY-2001; 2001AT-00000805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ24784 standard; DNA; 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lingnau K, Schellack C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ma.
Local S...
26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ24784;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                          strands
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ24784/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ð
                                                                                                                                                                                                                                                                The present invention relates to the isolation of a novel cytokine, alphanal Ligand and the polymucleotide encoding it. The invention also gives the sequence for the zalphal Ligand polymerotide stimulates proliferation of encoding it. The zalphal Ligand polypeptide stimulates proliferation of matural killer (NK) cells or NK cell progenitors, the activation of NK cells, proliferation of T-cells, proliferation of B-cells stimulated with anti-CD40 antibodies, stimulates an antigenic response in a mammal, and talphall Ligand polypeptide is also useful in preparing antibodies. The zalphall Ligand polypeptide is also useful in preparing antibodies that bind to zalphall Ligand epitopes. The zalphall Ligand epitopes or primers to clone regions of a zalphall Ligand gene, be used as probes or primers to clone regions of a zalphall Ligand gene, and in gene therapy. Zalphall Ligand may also be used to identify inhibitors of its activity, to enhance the generation of anti-tumour responses with or without the infusion of donor lymphocytes, and to activate or stimulate the immune system. The present sequence represents a sequencing primer used to sequence cibNA clones in the isolation of human zalphall Ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                 New zalphall ligand polypeptides and polynucleotides, useful for stimulating proliferation, activation, differentiation and/or induction of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      stranded polynucleotide tag; cleavage agent; gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
               Holly RD;
Hammond AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 26; DB 1; Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
               Foster DC,
Dillon SR, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .2e+02;
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0; Mismatches
               Sprecher CA,
Nelson AJ, I
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The present sequence is that of a thiosubstituted oligodeoxynucleic acid

Page 32; 57pp; English

Example 8;

Obtaining single stranded polynucleotide tags from a biological sample,

vaccine

New oligodeoxynucleic acid molecules useful for the preparation of

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Gaps

ö invention is based on the discovery that ODNs containing deoxyuridine invention is based on the discovery that ODNs containing deoxyuridine residues (U-ODNs) have an immunostimulatory effect comparable to, or in many instances greater than, ODNs containing CDG motifs, producing higher numbers of specific T cells to a given antigen. The U-ODNs do not induce CCG ODNs, are not dependent on a specific motif or a palindromic sequence. Use of a U-ODN for the preparation of a vaccine is claimed. Combining the U-ODN with an antigen strongly increases the potential of the antigen to raise the protection/immune response of a vaccinated individual. An example of the invention demonstrated the generation of a specific immune response edgins a vaccinated segmentic immune response against a malanoma-derived peptide (see es; PCR; zeig63; adhesin; salivary gland; dental carries; perinelial wound; urinary tract infection; vaginal infection; skin infection; primer; pro-inflammatory; chronic tissue damage; vascular system; diabetes; AIDS; lung infection; cystic fibrosis; lung dysfunction; digestive; salivary gland carcinoma; Pneumocystis carinii infection; emphysema; chronic bronchitis; prostate dysfunction; prostate adenocarcinoma; cell culture media; gene therapy; human chromosome 4q12-4q13; dentinogenesis imperfecta; dentin dysplasia type II. The invention relates to an isolated zsig63 polypeptide comprising at least 90% identity to an amino acid sequence which comprises domain 1 of Novel isolated zsig63 polypeptide, member of the adhesin family, useful for treating dental carries, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, Gaps ö 0.9%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 3.2e+02; 0; Indels Sequence 26 BP; 0 A; 0 C; 0 G; 1 T; 25 U; 0 Other; 100.0%; Prec. ... Human zsig63 PCR/sequencing primer ZC7764a. 2709 AAAAAAAAAAAAAAAAAAAAAAA 2734 26 AAAAAAAAAAAAAAAAAAAAA 1 Example 1; Page 29; 32pp; English. ABX93599 standard; DNA; 26 BP 03-AUG-2001; 2001US-00922469. 17-MAR-1999; 99US-0124820P. (first entry) Best Local Similarity 100. Matches 26; Conservative Adler DA, Sheppard PO; (ADLE/) ADLER D A. (SHEP/) SHEPPARD P O. WPI; 2003-328428/31. US2002173027-A1. skin infections 28-MAY-2003 21-NOV-2002. Synthetic. ABX93599; Query Match ABX93599/ ઠે 셤 

cc raig63, domain 2, domain 3, mature zaig63 and full length zaig63. Also included are the polymucleotide encoding zaig63, a zaig63 expression vector, a cultured call comprising the vector and expressing the protein, comprising the vector and expressing the protein, cc vector, a cultured call comprising the vector and expressing the protein, cc 126, 127-219 or 16-219 of zaig63 and an additional protein), using a mani-zaig63 antibody using zaig63 immunogenic peptides, Zaig63 is an anti-zaig63 antibody using zaig63 immunogenic peptides, Zaig63 is constituted to detecting in a test sample, the presence of antagonist of zaig63 protein activity. Zaig63 has antimicrobial activity and since constituted in salivary gland, can be used for treating dental carries, periodontal disease, thrush, and gastrointestinal content and protect against pathogenic colonization and other epithalial wounds. The polypeptides can be used to establish on the reating chronic, tissue damage particularly in areas having limited or damaged vascular system, e.g. in diabetes, and for treating chronic, tissue damage particularly in areas having limited or formunocompromised AlDS patients or in individuals that have undergone chemotherapy, radiation treatment, for treating lung infections e.g. in crystic fibrosis. Detection of zaig63 polypeptide at relatively high crystic fibrosis. Detection of zaig63 polypeptide at relatively high crystic fibrosis. Detection of zaig63 polypeptide at relatively high crystic fibrosis. Detection of zaig63 polypeptide at relatively high crystic fibrosis, prostete dysfunction. Zaig63 at also useful in dagnosing conditions associated with salivary gland or rung dysfunction in a useful in a dagnosis of effined cell colluture. The DNA is useful in a sectum that is commonly used in culture. The DNA is useful in age to relative and sequence is a primer used to isolate and sequence in undiaging type II). Zaig63 is an adhesin family media and sequence is a primer used to isolate and sequence in undiaging protects. ö Gaps ö ss; PCR; primer; antisense therapy; mRNA expression profile; 0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; ive 0; Mismatches 0; Indels Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other; 2708 TAAAAAAAAAAAAAAAAAAA 2733 26 талалалалалалалалалалалалал 1 BP. 07-SEP-2001; 2001US-00949305. 25-JAN-2000; 2000US-00494212 promoter containing primer. ACA62282 standard; DNA; 26 12-AUG-2003 (first entry) Query Match Best Local Similarity 100.0 Matches 26; Conservative Oligo (dT) primer #1. (EPIC-) EPICLONE INC. US2003022318-A1 Lin S, Ying S; 30-JAN-2003. Synthetic. ACA62282; RESULT 182 ACA62282, 셤 8

WPI; 2003-479488/45.

7 10:41:34 2006

Tue Nov

Improved polymerase thermocycling reaction for nucleic acid amplification, by thermal cycling of promoter-linked nucleic acid template synthesis and in vitro transcriptional amplification of nucleic acid sequences.

Example 4; Page 14; 28pp; English.

The invention relates to an improved polymerase thermocycling reaction (MI) for linear amplification of mucleic acid sequences, involves denatured (I) with a promoter-containing primer (PI), combining the denatured (I) with a promoter-containing primer (PI), a primer (P), a number of deoxymucleotide triphosphares and thouse of promoter-containing templates, contacting Pa with the denaturing the promoter-containing templates, contacting Pa with the denatured promoter-containing templates, containing templates, contacting Pa with the denatured promoter-containing demble-erranded mucleic acid templates are flanked by Pi in one on and Pi in the other and of the other orientation, transcribing the promoter-containing double-erranded moles transded mucleic as number of promoter-containing double-erranded DNA templates to form a number of containing double-erranded DNA templates to form a number of pulpares including the primer region of the promoter-containing double-erranded DNA templates, contacting the amplified RNA sequences, including the primer region of the promoter-containing double-erranded DNA templates, contacting the amplified RNA sequences, including the primer region of the promoter-containing double-erranded DNA templates, contacting the amplified RNA sequences, including the primer region of the promoter-containing double-erranded DNA templates, contacting the amplified RNA sequences of colls in both normal and diseased states, comparing the call should be generate multiple coffice by the calls in both normal and diseased states, emplifying the action of the calls in the normal and diseased state, determining the differences in mRNA composition of the call (S) in the diseased state, determining the malk and determining aberrant production, by the isolated mRNA will should be promoter protein producing an antisense cort the isolated mRNA and accompanies of the mRNA sequence of the mRNA in calling the companies of the mRNA and accompanies of the mRNA and using recompanies by the aberrant protein by dete produce second strand cDNA in the method of the invention

Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;

.; 0 0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; 0; Indels 100.0%; Pred. ... Local Similaricy hes 26; Conservative Query Match Best Loca Matches

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Gaps

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ADH44608 standard; DNA; 26 RESULT 183 ADH44608/c

(first entry) 25-MAR-2004 ADH44608; 2×4×2×

Human cDNA encoding Zalphall sequencing primer #2.

Human; 88; Zalphall ligand; Zalphall receptor; immune response; tumour progression; metastasis; tumour stasis; haematopoietic tumour; lymphoma; B cell tumour; systemic lupus erythematosus; rheumatoid arthritis; myssthemia gravis; diabetes; infectious disease; immunocompromised patient; HIV infection; vaccine; primer.

Ното варіепв.

US6605272-B2

12-AUG-2003

03-AUG-2001; 2001US-00923246.

99US-0123547P. 99US-0123904P. 99US-0142013P. 09-MAR-1999;

11-MAR-1999; 01-JUL-1999;

09-MAR-2000; 2000US-00522217

(ZYMO ) ZYMOGENETICS INC.

, Holly Hammond Foster DC, Dillon SR, F Sprecher CA, Nelson AJ, I Presnell SR, Johnston JV, Novak JE, Gross JA,

WPI; 2003-895283/82.

pathogen, useful for enhancing anti-tumor activity resulting in reduced tumor progression or metastasis, comprises administering zalphall ligand antigen or an Stimulating an immune response in a mammal exposed to polypeptide.

Example 7; SEQ ID NO 38; 103pp; English.

The invention fracts to stimulating an immunic response in a manufact comprising mature zalphall ligand polypeptide comprising a composition comprising mature zalphall ligand polypeptide comprising residues 32-162 comprising mature zalphall ligand polypeptide comprising residues 32-162 comprising; (a) determining (in) directly the level of antigen or pathogen present in the mammal; (b) administering a composition comprising salphall ligand polypeptide in a pharmaceutical vehicle; (c) determining (in) directly the level of antigen or pathogen in the mammal; (c) and (d) comparing the antigen or pathogen level in (a) with (b), where a change in the level indicates stimulation of immune response), and stimulating an immune response in a mammal exposed to an antigen or pathogen (comprising: (a) determining a level of immune response), and stimulating an immune response in a mammal exposed to an antigen or pathogen (comprising: (a) determining a level of antigen- or pathogen-comprising: (b) administering a composition comprising alphall ligand polypeptide in a pharmaceutical vehicle; (c) determining a post of c) comparing the level of the antibody in (a) with (b), where and increase in the antibody level indicates stimulation of immune response). The method is useful for stimulating an immune response in a mammal corresponse in a mammal antigen or pathogen, and for enhancing anti-tumour, a lymphoma or a B cell tumour may be a haematopoietic tumour, a lymphoma or a B cell tumour may be a haematopoietic tumour, a lymphoma or a B cell tumour may be a haematopoietic tumour, a lymphoma or a bealt of diseases arising from defects in the immune system; or systemic lupus erythematosus, rheumatoid arthritis, myasthemia gravis, or immunocymented por pathogen and diseases, treating and antigented in antigen or harlond antiful anticated and in immorymented and antigen and antigen and in antigen or pathogen. The invention relates to stimulating an immune response in a mammal immunocompromised patients, such as HIV+ patients and in improving vaccines. The present sequence is a sequencing primer used in the 

Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;

exemplification of the invention

Gaps ö 0.9%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 3.2e+02; 0; Indels 100.0%; Pred. No. ... 26; Conservative Best Local Similarity Query Match Matches

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2708 TAAAAAAAAAAAAAAAAAAAAA 2733

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US2004087526-A1.
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                                                                                     (LINS/) LIN S.
(JIHH/) JI H H.
                                                                                                                                                                              DNA-RNA hybrid
                                                                                                                  Ji HH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JUN-2004.
                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                               expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP19767;
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                   Lin S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 186
                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression; gene expredssion inhibition; eukaryotic cell characteristic; cell division rate; pigmentation; cancer; microbial infection; viral pathogenic infection; cancer cell proliferation; poly(dT)-26mer primer; ss; primer.
                                                                                                                             zalphall ligand; immunity; infectious disease; immunocompromised patient;
HIV; vaccine; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel isolated zalphall ligand polypeptide. The polypeptide of the invention may be useful for boosting immunity to infectious diseases and treating immunocompromised patients, such as HIV patients, as well as in improving vaccines. The current sequence is that of the PCR primer which was used in the exemplification of the invention.
                                                                                                          Sequencing primer SEQ 38 used to analyse human zalphall ligand clone DNA.
                                                                                                                                                                                                                                                                                                                             New zalphall ligand polypeptides, useful for boosting immunity to infectious diseases, and treating immunocompromised patients, such as human immunodeficiency virus (HIV) patients, or in improving vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene expression inhibition associated poly(dT)-26mer primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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100.0%; Pred. No. ...
0; Mismatches
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                                                 BP.
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                                                                                                                                                                                                                                         09-MAR-2000; 2000US-00522217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADO47862 standard; DNA; 26
                                                 ADI00944 standard; DNA; 26
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                                                                                      22-APR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                            (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                               Presnell SR,
Johnston JV,
                                                                                                                                                                                                                                                                                                           WPI; 2003-811003/76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 26; Conserv
                                                                                                                                                                               US2003125524-A1.
                                                                                                                                                            Homo sapiens
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                                                                     ADI00944;
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a targeted gene in a substrate, comprising a DNA-RNA hybrid. (I) is useful for inhibiting the expression of the targeted gene in a substrate; comprising a DNA-RNA hybrid. (I) is useful for inhibiting the expression of the targeted gene in a substrate. The substrate is a proxibation such as a viral or bacterial cell, or cukaryote is a mouse, rat, chimpanzee, preferably a human being. (I) is entaryote is a mouse, rat, chimpanzee, preferably a human being. (I) is caracteristic is chosen from expression of a protein, cell division rate and pigmentation. (I) has an effect that lasts at least three days. (I) is useful to inhibit the expression of messenger RNA in a cell. The messenger RNA is transcribed from a gene chosen from viral gene, oncogene, enzyme. (I) is useful for suppressing cancers, by knocking out cancer related genes, for preventing and treating microbial infections, preferably reducing viral pathogenic infection and for reducing the proliferation of cancer cells. This sequence represents a poly(dT)-26mer primer used in the creation of DNA-RNA hybrids for controlling gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention decribes a composition (I) for inhibiting the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; zalphall ligand; pharmaceutical; cancer; immune response; melanoma; tumour; solid tumour; haematopoietic tumour; lymphoma; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition useful for inhibiting the expression of a targeted gene in substrate, and for altering a characteristic of a eukaryote, comprises
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Pred. No. 3.2e+02;
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                                                                              12-NOV-2001; 2001US-0351183P.
18-JAN-2002; 2002US-00052486.
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19-MAR-2003; 2003US-00393450
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1es 26; Conservative
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26-FEB-2004; 2004US-00787442
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Best Local Similarity 100.
                                                                     Jimenez MC, Escobar IG,
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                       (GENO-) GENOMICA SAU.
(RUFF/) RUFFLES G K.
                                                                                                        WPI; 2004-507018/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2004260065-A1.
                                                                                                                                                                                              RNA hybrids.
                                                                                                                                                                                                                                                                                                                                                                                                         the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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(FOST/)
(HOLL/)
(GROS/)
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(PRES/)
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(DILL/)
(HAMM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 188
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                                                                                                                                                                                                                                                                                                    The invention describes an isolated polypeptide comprising a sequence of amino acid residues that is at least 90 or 95% identical to residues 41 (Gln) to 148 (Ile), or 32 (Gln) to 148 (Ile) of a sequence of 162 amino acids (SEQ ID NO:2, human zalphall ligand), fully defined in the specification. Also described are: a pharmaceutical composition comprising the polypeptide, and a vehicle; a method of trimulating an immune response in a mammal with melanoma; an method of stimulating an immune response in a mammal bearing a tumour; an isolated polymucleotide comprising a sequence of nucleotides contact an isolated polymucleotide comprising a sequence of nucleotides encodes a polypeptide that binds a receptor comprising 538 amino acids, fully defined in the specification; a pharmaceutical composition comprising the polymucleotide encoding, in a pharmaceutically acceptable comprising the polymucleotide encoding, in a pharmaceutically acceptable comprising the comprising parably linked a lamant. Sequence of nucleotides comprising the collowing parably linked a lamant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      elements a control element; and a DNA segment comprising the polynucleotide, and an isolated polynucleotide molecule comprising at least 10 nucleotides of the polynucleotide sequence of 642 bp, fully defined in the specification. The molecules, compositions and methods are useful for treating cancer, e.g. melanoma, solid tumour, haematopoietic tumour, or lymphoma. This sequence represents a primer used in the expression cloning of human cytokine zalphall ligand.
                                                                                                                                                                                                          useful for
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                                                                                                                                                                                                         New zalphall ligand polynucleotide and polypeptide molecules,
                                                                                                                      , Foster DC, Holly RD;
Dillon SR, Hammond AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                         Example 7; SEQ ID NO 38; 111pp; English.
                                                                                                                      Sprecher CA,
Nelson AJ, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAAAAAAAAAAAAAAAAAAAAA 1
99US-0123547P.
                                  01-JUL-1999; 99US-0142013P
09-MAR-2000; 2000US-00522217
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                                                                                    (ZYMO ) ZYMOGENETICS INC
                                                                                                                    Presnell SR,
Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Da(26) biotin primer.
                                                                                                                                                                      WPI; 2004-440401/41.
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les 26; Conserv
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09-MAR-1999;
11-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUN-2004
                                                                                                                  Novak JE,
                                                                                                                                      Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADQ80457;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Matches
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The present invention relates to experimentally analysing boundaries within polymeric DNA or RNA molecules comprises hybridizing first and second different RNA molecules, derived from first and second samples, to provide RNA / RNA hybrids. The product of the method above is used as a probe. The method is useful in experimentally analysing boundaries within polymeric DNA or RNA molecules. The methods are useful in analysing polymeric nucleic acid sequence variations and in identifying molecules of therapeutic interest. The present sequence represents a RNA primer of
                                                                                                                                  Experimentally analyzing boundaries within polymeric DNA or RNA molecules, useful in analyzing polymeric nucleic acid sequence variations, comprises hybridizing different RNA molecules to provide RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stem cell; cell culture; PCR; primer; ss; zalphall ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 26; DB 1; Length 26;
Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
Cimadevilla JCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human zalphall ligand-specific PCR primer - SEQ ID 38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 14; 36pp; English
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Gallego SC,
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11-WAR-1999; 99US-0123904P.
01-JUL-1999; 99US-0142013P.
09-WAR-2000; 2000US-00522217.
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Synthetic
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                                                                                                                                                                                                                                                                                                                 RESULT 190
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                                                                                                                        The invention comprises a fusion protein that contains a zalphall ligand and a cytokine polypeptide (e.g. IL-2, IL-4, IL-15 or GM-CSF), the fusion protein of the invention binds to the human receptor protein. The protein of the invention is useful for stimulating the proliferation and/or development of hematopoietic cells. The protein of the invention is also useful in autologous marrow culture. The present DNA sequence represents a PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dental carries; candida infection; fungicide; infection; periodontal disease; antiinflammatory; mouth disease; gastrointestinal disease; gastrointestinal disease; gastrointestinal disease; female gential tract infection; antimicrobial; uropathic; genitourinary disease; female genital tract infection; antimicrobial; gynecology and obstetrics; skin infection; dermatological disease, wound healing; vulnery; injury; acquired immune deficiency syndrome; anti-hiv; immune disorder; cancer; cytostatic; neoplasm; lung infection; antimicrobial; respiratory-gen; respiratory disease; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zaig63; microbial infection; tooth disease; antibacterial; mouth disease;
                                                New zalpha 11 Ligand fusion protein, useful for stimulating the proliferation and/or development of hematopoietic cells in vitro and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide (I) encoding a zaig63 polypeptide, useful for diagnosing and treating microbial infections, e.g. dental carries, thrush, gastrointestinal disease, skin infection, or epithelial wounds, AIDS, lung infections, or cancer.
                                                                                                                                                                                                                                                                            Gaps
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  Hammond AK;
                                                                                                                                                                                                                                                  Query Match 0.9%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 3.2e+02; Matches 26; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Zsig63 cDNA cloning and sequencing primer, ZC7764a.
                                                                                                                                                                                                                             Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
 Dillon SR,
                                                                                                                                                                                                                                                                                                     2708 TAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                 Example 7; SEQ ID NO 38; 110pp; English.
                                                                             vivo, and in autologous marrow culture.
                                                                                                                                                                                                                                                                                                                    26 TAAAAAAAAAAAAAAAAAAAAA 1
  Nelson AJ,
                                                                                                                                                                                                                                                                                                                                                                                           ADY96657 standard; DNA; 26 BP
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03-AUG-2001; 2001US-00923236.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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  Johnston JV,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2005-241320/25.
                        WPI; 2005-038783/04
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                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUN-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer; 88
  Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                    ADY96657;
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                                                  The present invention relates to zsig63, a novel secreted salivary protein and its encoding polynucleotide. Zsig63 is a member of the adhesin family. The invention is useful for diagnosing and treating microbial infections such as dental carries, periodontal disease, thrush, gastrointestinal disease, unimary tract infection, vaginal infection, skin infection or epithelial wounds and lung disfunctions such as AIDS, lung infections and cancer. The invention is also useful in gene therapy. The present sequence is human Zsig63 cDNA cloning and sequencing primer. This sequence to obtain the full-length Zsig63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel method for parallel sequence analysis of nucleic acids (NA) by optical means using a novel solid phase (SP). The SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids, so the background remains low even after prolonged an repeated contact of the solid phase with high concentrations of labeled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
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100.0%; Pred. No. 3.2e+02;
tive 0; Mismatches 0;
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Example 1; SEQ ID NO 7; 33pp; English.
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/note= "OTHER= Cy3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2006-040183/05.
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Best Local Similarity
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Best Local Similarity 100.
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GENOVOXX GMBH.
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                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                      Synthetic.
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                                              RESULT 192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parallel sequencing of nucleic acids by optical methods, by cyclic primer -matrix extension, using a solid phase with reduced non-specific binding of labeled components.
 reagents. The present sequence is that of an oligonucleotide which was used in the development of the novel method of the invention.
                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                              DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
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0
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                                                                    0.9%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 3.2e+02;
ive 0; Mismatches 0; Indels
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                                         .Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                               Location/Qualifiers
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/note= "OTHER= Cy3"
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                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                   26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ′*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CHEN/) CHERKASOV D.
(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
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                                                                                    Similarity
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Matches 26; Conserv
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The
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modified_base
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                                                                       Query Match
                                                                                                                                                                                                                                                            AEE86828;
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                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HENN/)
(GENO-)
                                                                                       Best Loca
Matches
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This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid phase of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                            DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing;
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                                                                                                                                            Novel solid phase-related oligonucleotide Oligo dA26-Cy3 #2
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.00.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                              /note= "OTHER= Cy3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 62; 88pp; German
                                                                                                                                                                                                                                                                                                                                                                       OTHER
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/mod_base=
AEE86844 standard; DNA; 26
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2709 AAAAAAAAAAAAAAAAAAAAA 2734

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26-MAY-2004; 2004DE-10025744. 26-MAY-2004; 2004DE-10025744.

Page 100

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Surface of a solid support, useful for multiple parallel analysis nucleic acids by optical methods, having low non-specific binding labeled components.

Hennig C;

Cherkasov D,

WPI; 2006-081126/09.

(GENO-) GENOVOXX GMBH. CHER/) CHERKASOV D.

HENNIG

HENN/)

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                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel surface of a solid phase (SP), useful in methods for parallel analysis of many individual nucleic acids (NA) by optical methods. The novel SP is useful for multiple parallel sequencing of nucleic acids and shows reduced non-specific binding of labeled or unlabeled nucleotides and nucleic acids. The present sequence is that of an oligonucleotide which was used in the development of the novel solid
                                                                                                                                                                                                                                                                                                               Surface of solid phase, useful for parallel, optical analysis of many nucleic acids, has reduced non-specific binding of labeled components
DNA chip; biochip; DNA detection; DNA microarray; DNA sequencing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "labelled with Cy3"
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                                             Location/Qualifiers
                                                                                /mod_base= OTHER
/note= "OTHER= Cy3"
                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 62; 88pp; German.
                                                                                                                                                                   26-MAY-2004; 2004DE-10025745
                                                                                                                                                                                          26-MAY-2004; 2004DE-10025745
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide dA26-Cy3.
                                                                                                                                                                                                                                                                  Cherkasov D, Hennig C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     phase of the invention.
                                                                      *tag=
                                                                                                                                                                                                                 (HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
(CHER/) CHERKASOV D.
                                                                                                                                                                                                                                                                                         WPI; 2006-040182/05.
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Best Local Similarity
Matches 26; Conserv
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                                                                                                                     DE102004025745-A1.
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modified_base
                                                          modified base
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                       Synthetic
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This invention describes a novel solid phase in which the surface shows invention also describes, a) a solid phase in which the surface shows reduced non-specific binding of ladd components; b) methods for reduced non-specific binding of ladd components; b) methods for reduced non-specific binding of ladd components; b) methods for preparing the solid support and c) methods of parallel analysis of many nucleic acid by optical methods, using the solid support. The solid phase is preferably part of a device that through a linker. The solid phase is preferably part of a device that allows fluid exchange and it is permeable to light in the wavelength cregions 200-400; 200-2000 or 400-800 nm. An external layer of solid cupport is removed, then the nucleic acid is coupled to it, optionally cupport is removed, then the nucleic acid is coupled to it, optionally cupport is removed, then the nucleic acid is coupled to it, optionally a feer attachment of a linker layer. Alternatively, after removing the extends of specifically phosphate, sulfate or carboxy.

Containing monomers or polymers) can be coupled to the surface, after coupling, optionally by a chemical reaction with hydrofluoric acid or containing monomers or polymers) can be completed to the surface is attachment or synthesis of nucleic acide. Only part of the surface is containing monomers or polymers) can removed particularly by a chemical reaction with hydrofluoric acid or analyzed represent a single population or many different populations and all subsequent steps are done in a liquid phase. The nucleic acid or on any population or an any serious sees is involved or linear polymer; (archidate promote or linear polymer; (archidate parallel analysis uses components labeled with ribo. A deoxyribonucleoside triphosphates, in which the label is cleavable. Particularly analysis involves cyclic sequencing and a preferred method contains the nucleoside triphosphate; for the solid support, with formation of a cetensible primer-matrix complex; performing acid to the s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotides or nucleic acids,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gapa
                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel solid support surface for parallel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   so assay sensitivity is improved. This sequence represents an oligonucleotide used to illustrate the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     parallel sequencing of nucleic acids. The surface specific binding of labeled and unlabeled nucleoti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.0%; Pred. No. 3.2 (es 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2734
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                                                                                                                                                                                                                                                                                                                                                                   Disclosure, Page 62; 88pp; German.
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Optical DNA analysis process-related oligonucleotide dA26-Cy3 #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faget, more efficient and cheaper than prior at. Further claimed is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dA26-Cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                      Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                   ss; dna detection; DNA sequencing; DNA amplification; oligo dA26-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                            Optical DNA analysis process-related oligonucleotide dA26-Cy3 #1.
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100.0%; Pred. No. 3.2e+02;
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Conservative 0; Mismatches
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                                                                                                                                                                                  /*tag= a
/mod_base= 5'-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 66; 94pp; German.
          AEF94771 standard; DNA; 26 BP.
                                                                                                                                                                                                                                                                 26-MAY-2004; 2004DE-10025695
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                                                     20-APR-2006 (first entry)
                                                                                                                                                                                                                                                                                                                                                           Cherkasov D, Hennig C;
                                                                                                                                                                                                                                                                                                                         HENNIG C.
GENÓVOXX GMBH.
                                                                                                                                                                                                                                                                                                             (CHER/) CHERKASOV D.
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Best Local Similarity
Matches 26; Conserv
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                                                                                                                                                           Key
modified_base
                                                                                                                         Unidentified
                                                                                                                                                                                                                                          23-FEB-2006
                                                                                                                                     Synthetic
                                AEF94771;
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(GENO-)
AEF94771
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dA36-Cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Optical fluorescent parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
ss; dna detection; DNA sequencing; DNA amplification; oligo dA26-Cy3
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100.0%; Pred. No. 3.2e+02;
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                                                                                                                                Location/Qualifiers
                                                                                                                                                                                /*tag= b
/mod_base= 5'-Cy3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hennig C;
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(GENO-) GENOVOXX GMBH.
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                                                                                                                             Key
modified_base
                                                  Unidentified
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                                                                             Synthetic.
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AEF94785 standard; DNA; 26

RESULT 196

**AEF94785** 

AEF94785;

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20-APR-2006 (first entry)

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AEF94730;
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                                                                                                                                                                                                                                                                    This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. The present sequence is that of oligonucleotide da26-cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                     Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88; dna detection; DNA sequencing; DNA amplification; oligo dA26-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Optical DNA analysis process-related oligonucleotide dA26-Cy3 #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%; Score 26; DB 1; Length 26; 00.0%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

10.0%; Jocal Similarity 100.0%; Pred. No. 3.2

Matches 26; Conservative 0; Mismatches
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 Location/Qualifiers
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/mod_base= 5'-Cy3
                   /*tag= b
/mod_base= 5'-Cy3
                                                                                                                                                                                                                                                     Example 2; Page 68; 95pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEF94755 standard; DNA; 26 BP
                                                                                              26-MAY-2004; 2004DE-10025694
                                                                                                                26-MAY-2004; 2004DE-10025694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                              Cherkasov D, Hennig C;
                                                                                                                                     (CHER/) CHERKASOV D.
(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                  WPI; 2006-185818/20.
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                                                     DE102004025694-A1
  Key
modified_base
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
Synthetic.
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                                                                         23-FEB-2006
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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction, followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the sequence repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is sequence the interpretation of oligonalectide dA26-Cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                         Optical fluorescent ultra-high parallel process to analyse nucleic acid chains in which a sample solid is bound with a primer-matrix complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; dna detection; DNA sequencing; DNA amplification; oligo dA26-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Optical DNA analysis process-related oligonucleotide dA26-Cy3 #2.
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Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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/mod_base= 5'-Cy3
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 5; Page 67; 95pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEF94730 standard; DNA; 26 BP
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26-MAY-2004; 2004DE-10025694.
                                                             26-MAY-2004; 2004DE-10025694
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                                                                                                                                                                                                                                             Cherkasov D, Hennig C;
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(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                     (CHER/) CHERKASOV D.
(HENN/) HENNIG C.
(GENO-) GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                       WPI; 2006-185818/20.
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modified_base
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Tue Nov

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This invention relates to a novel optical fluorescent process to analyse nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction. Marked nucleotides washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their signals removed. The signals removed. The solid phase is then washed and the signals repeated as necessary. The Nucleic acid chain sequence is then reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide dA26-Cy3 which was used in
                                                                             Ultra-high parallel analysis process to analyse nucleic acid chains in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ultra-high parallel analysis process to analyse nucleic acid chains in which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; dna detection; DNA sequencing; DNA amplification; oligo dA26-Cy3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Optical DNA analysis process-related oligonucleotide dA26-Cy3 #1.
                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 3.2e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                              development of the novel process of the invention.
                                                                                            which a sample solid is bound and substrate material.
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                  Baeuml E;
                                                                                                                            Example 2; Page 95; 141pp; German.
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Matches 26; Conservative
                  Hennig C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cherkasov D, Hennig C,
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GENOVOXX GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHER/) CHERKASOV D.
                                                WPI; 2006-185820/20.
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                Cherkasov D,
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Gaps

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                                                                nucleic acid chains. Using the method, a sample solid is bound with a primer-matrix complex resulting in a cyclic reaction. Marked nucleotides are incorporated in the primer matrix by enzyme reaction. followed by washing of the solid phase. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The marked nucleotides are detected and their co-ordinates logged and the signals removed. The solid phase is then washed and the reconstructed using the signals. The process is faster, more efficient and cheaper than prior art. Further claimed is that the process able to determine many sequences in parallel. The present sequence is that of oligonucleotide da26-cy3 which was used in the development of the novel process of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preparation of normalised, subdivided cDNA libraries from mRNA - by reverse transcription and amplification, used to screen for new genes and interacting proteins, potential drugs, and for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to preparation of a normalised, subdivided library of amplified cDNA from the coding regions of mRNA in a sample. The method involves reverse transcription, with at least one cDNA primer of formula 5'-Conl-dTn2-Vn3-Nn4 to form first stand cDNA where Conl = any sequence of 1-100 nucleotides; dT = deoxythymidinyl; n2 is at least 1; n3 and n4 are both 0, or n3 is 1 and n4 is at least 1; followed by second strand cDNA synthesis using the first strand as template and a second cDNA
                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Normalised, cDNA library, mRNA cloning, reverse transcription, immobilise, screening, hybridisation, nucleic acid amplification, expression pattern, drug development; PCR primer, RT-PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                               0.9%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                   invention relates to a novel optical fluorescent
                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 26 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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               Example 5; Page 95; 141pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 29; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anchored poly T RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                  26; Conservative
                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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27-MAR-1998;
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ID AAV7
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The linear multimer was produced by rolling circle synthesis in an example of the method of the invention for synthesising an RNA oligonucleotide, comprising combining a single-stranded circular coligonucleotide template comprising at least one copy of a nucleotide sequence complementary to the sequence of the desired RNA oligonucleotide with at least 2 types of ribonucleotide triphosphate and a polymerase enzyme to yield a single-stranded RNA oligonucleotide multimer complementary to the circular oligonucleotide template, where the RNA oligonucleotide multimer compressed multiple copies of the desired RNA oligonucleotide. The methods can be used for producing RNA oligonucleotides having a specific sequence and well defined ends. The structions produced can be used as probes, standards and diagnostic or therapeutic appears. They can be used for modifying the structure or function of a target molecule. They can also be used to
                                                                                                                                                                                                                                                                                                                                                                                         ö
primer of a similar formula, in the presence of DNA polymerase I (or its Klenow fragment) and amplification of double-stranded cDNA with a set of amplification primers. Comparison of cDNA in the prepared library with a database (a computer-generated list of molecular weights of restricted DNA fragments of known sequence) is used to determine presence of an expressed protein in a cell, also to detect changes in such expression (particularly for diagnosis of disease). Surfaces (chip) having amplified cDNA stably immobilised on it, obtained by a similar method, are used to screen for genes of a particular family, by hybridisation with nucleic acid from the family (to identify new genes) and to detect differences in expression patterns between cells. The polypeptides expressed by the libraries can be used for drug development. Sequences AAV71935 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    вв; RNA oligonucleotide; probe; standard; diagnostic; therapeutic agent.
                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthesis of oligo:nucleotide(s) - using a single-stranded circular oligo:nucleotide template ribonucleotide tri:phosphate(s) and a polymerase to form multimer(s) which can be cleaved.
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                                                                                                                                                                                                                                                                                                                                                DB 1; Length 27; 3.3e+02;
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                                                                                                                                                                                                                                                                                                      Sequence 27 BP; 2 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100.0%; Pred. No. 3.3 nes 26; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 26;
100.0%; Pred. No.
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oligonucleotide, comprising combining a symmetraling unit interior.

Oligonucleotide, comprising combining a symmetral controlled to oligonucleotide template with at least two types of ribonucleotide complate with at least two types of ribonucleotide coligonucleotide template with at least two types of ribonucleotide multimer complementary to the circular oligonucleotide multimer complementary to the circular oligonucleotide multimer comprises multiple template, where the RNA oligonucleotide The method is useful for synthesising an RNA oligonucleotide with well-defined ends. The circular oligonucleotide is introduced into the cell using direct injection, calcium phosphate treatment, lipid-mediated delivery, or cation-mediated delivery. The method is useful for treating bacterial and/or viral infections in mammals, particularly drug resistant infections in mammals, particularly drug resistant infections, and for producing double stranded DNA oligomers. The method is performed in the absence of an oligonucleotide primer, or without the addition of auxiliary proteins. This sequence represents an oligonucleotide used in the method of the invention.
                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA oligonucleotide synthesis; ribonucleotide triphosphate; polymerase; electroporation; calcium phosphate treatment; lipid-mediated delivery; cation-mediated delivery; bacterial infection; viral infection; drug resistant infection; double stranded DNA oligomer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthesizing RNA oligonucleotide involves combining single-stranded circular oligonucleotide, ribonucleotide triphosphate and polymerase enzyme to yield desired RNA complementary to circular oligonucleotide
                                                                                                             Gaps
                                                                                                             .;
0
                                                                     Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method for synthesising an RNA
                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29 BP; 0 A; 0 C; 2 G; 27 T; 0 U; 0 Other;
                               Sequence 29 BP; 0 A; 0 C; 2 G; 27 T; 0 U; 0 Other;
                                                                       0.9%; Score 25.8; DB 1; 93.1%; Pred. No. 3.5e+02; ative 0; Mismatches 2;
cleave disease-associated RNA, DNA or protein
                                                                                                                                                  2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2737
                                                                                                                                                                                      29 AAAAAAAAAACAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; SEQ ID NO 6; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93US-00047860.
95US-00393439.
97US-00805631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-2001; 2001US-00997931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-1997; 97US-00805631.
11-MAY-2000; 2000US-00569344.
                                                                                                                                                                                                                                                                                       ADC65873 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                      DNA oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYRP ) UNIV ROCHESTER
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                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003087241-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-APR-1993;
23-FEB-1995;
                                                                                 Ma.
Local St...
27;
                                                                                                                                                                                                                                                                                                                                                                 18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                             ADC65873;
                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kool ET;
                                                                                                   Best Loca
Matches
                                                                                                                                                                                                                                                    RESULT 203
                                                                                                                                                                                                                                                                       ADC65873/
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29 AAAAAAAAAAAAAAAAAGAAAGAAAA 1

(first entry)

29-JUL-2004

ADO81069;

ADO81069 standard; DNA;

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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA semple containing at least one segment of (1); and cast emplate a DNA semple containing at least one segment of (1); and cast emplained of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or predisposition to a disease, associated with a gene that includes one or include PML; and prediagnosis (M3) of diseases associated with gene that includes one or more PML; and prediagnosis (M3) of diseases, especially markers, in a disease related gene, that are associated with a predisposition to diseases and for prediagnosis of such diseases, especially prion diseases to be also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length.of amplicons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        region of the cow prion
                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease predisposition, microsatellite marker, prion disease, cystic fibrosis, malignant hyperthermia syndrome; metabolic disease, milk protein, hormone; transcription factor; pT7-blue-vector; cow;
                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence represents a primer used to genotype a region of the protein (PrP) comprising a polymorphic microsatellite locus.
                  Length 29;
                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cow prion protein microsatellite locus primer #77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene typing; polymorphic microsatellite loci; PML;
          Score 25.8; DB 1;
Pred. No. 3.5e+02;
0; Mismatches 2;
                                                                                                                                                  2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2737
                                                                                                                                                                                          29 AAAAAAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                               ADO81065 standard; DNA; 29 BP
              0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2002; 2002DE-01036711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JUL-2004 (first entry)
                                                                                   27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYHO-) UNIV HOHENHEIM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-215730/21.
                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DE10236711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Geldermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30s taurus
Query Match
Best Local S
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                                                                            Matches
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The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and castermining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease. Associated with a gene that includes one or predisposition to a disease, associated with a predisposition to a disease. Associated with a predisposition to consider that a seasociated with a predisposition to diseases associated with a predisposition to diseases associated with a predisposition to diseases and for predisposis of such diseases, especially prion diseases to the also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, metabolic diseases also to type genes that encode milk proteins, particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion comprising a polymorphic microsatellite locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                         disease predisposition, microsatellite marker, prion disease, cystic fibrosis, malignant hyperthermia syndrome, metabolic disease, milk protein, hormone, transcription factor, p77-blue-vector, cow,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 29 BP; 0 A; 2 C; 0 G; 27 T; 0 U; 0 Other;
                                                                               Cow prion protein microsatellite locus primer #81.
                                                                                                                        gene typing; polymorphic microsatellite loci; PML
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25.8; DE
Pred. No. 3.5e+
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAAAAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 28; 64pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Han Y;
                                                                                                                                                                                                     microsatellite; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                               09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                                                                                       09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 93.1%;
les 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Geldermann H, Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYHO-) UNIV HOHENHEIM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-215730/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         length of amplicons
                                                                                                                                                                                                                                                                                  DE10236711-A1.
                                                                                                                                                                                                                                                                                                                          26-FEB-2004.
                                                                                                                                                                                                                                             30s taurus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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AAQ83940 standard; DNA; 30 BP. RESULT 206 AAQ83940 П

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Gaps

; 0

0.9%; Score 25.8; DB 1; Length 29; 13.1%; Pred. No. 3.5e+02; ve 0; Mismatches 2; Indels

Query Match 0.9%; Best Local Similarity 93.1%; Matches 27; Conservative

Sequence 29 BP; 0 A; 2 C; 0 G; 27 T; 0 U; 0 Other;

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2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2737

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The sequences given in AAQ83918, AAQ83952 and AAQ83940 are used in the construction of an oligonucleotide clamp. The clamp is a comb-type branched polymer which has 3' termini and was used to bind a target bequence comprising a segment of the HIV pol and nef genes in single stranded or double stranded forms. An oligonucleotide clamp is a compound complex after specifically binding to the target polynucleotide.

Complex after specifically binding to the target polynucleotide or a stable circular complex after specific binding to the target molecule and one or more pairs of binding molecies cone or more oligonucleotide moieties capable of specific binding to the target molecule and one or polynucleotide, the binding moieties of a pair are bought into inxapposition so that they form a stable covalent or non-covalent linkage or complex. The interaction of the binding moieties effectively clamps the specifically annealed oligonucleotide moieties to the target the specifically annealed oligonucleotide moieties to the target polynucleotide. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthesis of branched polymers and novel branched polymeric structures used as molecular probes esp. for detecting poly-nucleotide (s)\,\,.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                HIV; pol; nef; oligonucleotide clamp; branched; macromolecule; ss.
                                                                                                Oligonucleotide clamp o, for producing comb-type brached polymer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.9%; Score 25.8; DB 1; Length 30; Best Local Similarity 93.1%; Pred. No. 3.6e+02; Matches 27; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                            /*tag= a
/note= "Modified with SP(O-)(=O)-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 27 A; 3 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2706 ACTAAAAAAAAAAAAAAAAAAAAAAAA 2734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 ACACAAAAAAAAAAAAAAAAAAAAAA 30
                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Page 33; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                (LYNX-) LYNX THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                              93US-00087386.
                                                                                                                                                                                                                                                                                                                                                            94WO-US007557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF60462 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide clamp #22
                                                                                                                                                                                                                                          *tag=
                                                  (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-060944/08.
                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                                                            05-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                02-JUL-1993;
                                                                                                                                                                                                                                                                                            W09501365-A1
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                                              25-MAR-2003
04-OCT-1995
                                                                                                                                                                                                                                                                                                                            12-JAN-1995
                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF60462;
                AAQ83940;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF60462
0 X 2 X 5 X 8
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The present invention relates to a method for synthesising a branched or multiply connected macromolecular structure, comprising oligonucleotide clamps (OC). The macromolecular structure is capable of specifically binding to a target molecule, and can therefore be used as probes. At least one OC comprises a target binding sequence that binds specifically and stably with the target molecule, and at least two OCs comprises signal generation moieties capable of generating a detectable signal in the presence of the target molecule. In addition the OCs are connected to one another by thioalkylamino, or thiophosphorylalkylamino bridges. The
                                                                                                                                                                                                                                                                           Synthesizing branched nucleic acids useful as diagnostic and molecular probes, involves combining first units having haloalkylamino groups and second units having thiol or phosphorothioate groups.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genotype; plant; rice; semi-dwarf; sd-1; polymorphism; detection;
characteristic; single nucleotide polymorphism; SNP; genotyping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence is an OC used in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 27 A; 3 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rice semi-dwarf (8d-1) DNA fragment SEQ ID NO:26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 25.8; DB 1;
Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2706 ACTAAAAAAAAAAAAAAAAAAAAAAAAAA 2734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                       Example 8; Col 19; 20pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%;
Best Local Similarity 93.1%;
Matches 27; Conservative 0
                                                                                                                               97US-00787321.
                                                                                                                                                           96US-0009918P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
          Oligonucleotide clamp; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chromosome 1; gene; ds
                                                                                                                                                                                                                                                  WPI; 2001-201911/20.
                                                                                                                                                                                         CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003070934-A1.
                                                                                                                                                                                        (FARB ) BAYER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oryza sativa.
                                         Unidentified.
                                                                                                                             03-JAN-1997;
                                                                                                                                                            12-JAN-1996;
                                                                     US6180777-B1
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                                                                                                   30-JAN-2001
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ADA26181/c
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Gaps

7 10:41:34 2006

Tue Nov

Suzuki

Yoshino R,

Kitazawa N,

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Disclosure; Page 15; 104pp; Japanese.
      WPI; 2003-697617/66.
                                                                                                     25-MAR-2003
29-NOV-1995
                                                                                                                Normalised
  Minobe Y,
                                                                                                 AAQ87894;
                                                                      Query Match
                                                                                         RESULT 209
                                                                                           AAQ87894/
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                                                                                  셤
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셤 ઠે The present invention describes a method for judging the genotype of a region around a plant semi-dwarf (sd-1) gene in which polymorphisms are present, by detecting the polymorphisms. Also described: (1) examining semi-dwarf characteristics of a plant using the judgment method with detection of polymorphisms; (2) oligonucleotides for amplifying sd-1 DNA regions, which are primers for judging the genotype of a region around a plant sd-1 gene; (3) reagents for judging the genotype of a region around a plant sd-1 gene; (3) reagents for judging the genotype of a region around a plant sd-1 gene; (4) reagents for judging the genotype of a region around a plant sd-1 gene; (4) reagents for judging the genotype of a region around a plant sd-1 gene; which is applicable in genotyping by (3) CAPS ((derived) cleaved amplified polymorphic sequence) for examination of the semi-dwarf character of rice to identify desirable strains e.g. with high crop yield, peet realstance and resistence to flooded water. The method is easy and quick, in which a seedling is required for studying single nucleotide polymorphisms (SNPs) for genotyping, without needing cultivation of seedling to fully-grown plant for judging heterozygote and distinguishing morphology. The present sequence represents a rice sd-1 invention. Rice sd-1 is located on chromosome 1. ; 0 cDNA library; directionally cloned cDNA library; screening; Gaps .; 0 3; DB 1; Length 30; 3.6e+02; 2; Indels Normalised library first strand cDNA synthesis primer. Sequence 30 BP; 0 A; 3 C; 0 G; 27 T; 0 U; 0 Other; 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAA 2737 0; Mismatches 0.9%; Score 25.8; 93.1%; Pred. No. 3. 30 AAAAAAAAAAAAAAAAAAAAA 2 Location/Qualifiers AAQ87894 standard; DNA; 32 (first entry) Best Local Similarity 93.1 Matches 27; Conservative (revised) hybridisation; ss. misc\_feature Synthetic.

(UYCO ) UNIV COLUMBIA NEW YORK.

94WO-US010821. 93US-00126594.

23-SEP-1994;

24-SEP-1993;

WO9508647-A1

30-MAR-1995.

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Human tissues were obtained for construction of a variety of cDNA libraries, including infant brain, adult brain and adult hippocampus. Bach of the cDNA libraries had a characteristic sequence identifier, provided by the oligonucleotide utilised to prime first strand cDNA synthesis (see AAQ87894-Q87907 for these primer sequences; all these synthesis (see AAQ87894-Q87907 for these primer sequences; all these forfiners have the PacI restriction site for directional cloning of cDNAB). Each of the libraries was propagated in the form of single-stranded (ss) circles and normalised separately by a novel method. The method comprises: generating fragments complementary to the 3' non-coding sequence of the special duplexes; melting and reassociating them to appropriate Cot; and purifying the unassociated so circles to generate a normalised cDNA library. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying a candidate polymorphic repeat within a coding sequence, for understanding or treating genetic disease, comprises detecting tandem repeats in a target coding sequence and scoring the repeats for polymorphic probability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPOUS; Rep.Y; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Predreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                              New normalised directional cDNA libraries - used for isolating novel cDNA's, including tissue-specific and development-specific DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.9%; Score 25.6; DB 1; Length 32; 87.5%; Pred. No. 3.8e+02; Live 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 32 BP; 4 A; 0 C; 0 G; 28 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST polymorphic DNA repeat polynucleotide #153.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 ААААААААААААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAATTAATTAAAAAAA 1
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                                                                                                                                                        Disclosure; Page 45; 186pp; English.
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                  Efstratiadis A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28; Conservative
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                                                         WPI; 1995-139615/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                    Soares MB,
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Best Local &
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Judging the genotype of a region around a plant sd-1 gene with polymorphism-obtained markers isolated by positional cloning, useful in genotyping for examination of semi-dwarf character of rice.
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/note= "characteristic sequence identifier"
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                                                       repeat within a coding sequence (expressed sequence teg, EST), which repeat within a coding sequence (expressed sequence teg, EST), which the repeats for polymorphic probability and generating a dataset correlating the repeat with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                snake venom gland RNA. Polynucleotides encoding snake venom protease (SVP) were isolated from brown snake ADG83825, coastal taipan ADG83827, inland taipan ADG83829, red belly ADG83831, tiger ADG83833 and rough scale ADG83835 snake venom gland cDNA libraries. The invention is based on the discovery of the cofactor-independent prothrombin activating activity of these SVPs. The SVPs can be used e.g. to promote haemostasis
                                     invention discloses a method for identifying a candidate polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New snake venom protease preparation useful in promoting hemostasis and preventing blood loss such as during surgery, in treating wounds resulting from accidents and other types of injury or trauma, or as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present sequence is that of a primer used for cDNA synthesis from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Snake venom; protease; prothrombin activator; haemostatic; vulnerary;
fibrin glue; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.9%; Score 25.4; DB 1; Length 27; Best Local Similarity 96.3%; Pred. No. 3.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                      Sequence 27 BP; 1 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lavin M;
Example; Col 717; 588pp; English
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07-MAR-2003; 2003AU-00901033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          surgical sealant or adhesive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-APR-2003; 2003WO-AU000406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG83852 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer for cDNA synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYOU ) UNIV QUEENSLAND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a polynucleotide derived from the 4q12-4q13 region of human chromosome 4 and encoding a zsig63 polypeptide, a secreted salivary protein with anti-microbial activity. Due to their microbial activity, the sequences can be used in the study of microbial infections, e.g. for recombinant production of anti-microbial proteins. The sequences can be used in the treatment of tooth decay, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, skin infections, epithelial wounds, chronic tissue damage, acquired immunodeficiency syndrome (AIDS), cyptic fibrosis, lung infections, sarcoidosis, emphysema and chronic bronchitis. This sequence represents a sequencing primer for cDNA encoding human zsig63
and prevent blood loss such as during surgery or for treatment of wound resulting from injury or trauma, and may be useful as a topical fibrin 'glue' or 'sealant'.
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                    0; Indels
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                                                                                                                                                       Sequence 27 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 2 Other;
                                                                                                                                                                                                                0.9%; Score 25.4; DB 1;
92.6%; Pred. No. 3.6e+02;
tive 2; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human zsig63 cDNA sequencing primer ZC7231.
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                                                                                                                                                                                                                                                                                                                                                       2707 CTAAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                                                                                                                            Best Local Similarity 92.6
Matches 25; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Adler DA, Sheppard PO;
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                                                                                                                                                                                                                              Query Match
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RESULT 213

## BAAAAAAAAAAAAAAAAAAAA 1

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ABS52637 standard; DNA; 26 BP
                                                                                    15-NOV-2002 (first entry)
                                                  ABS52637;
ABS52637
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Human; secreted salivary protein; zsig63; immunogen; zsig63-cytokine; antibody-cytokine; in vivo killing; pathological microbe; bacteria; fungal; viral; infection; salivary gland; anti-microbial; dental caries; tooth decay; periodontal disease; thrush; gastrointestinal disease; urinary tract infection; vaginal infection; kin infection; microficar; epithalial wound; pathogenic colonisation; invasion; pro-inflammatory; chronic tissue danage, vascular system; diabetes; anti-inflammatory; incompetent immune system; AIDS; acquired immunodeficiency syndrome; chemotherapy, radiation treatment; lung infection; cystic fibrosis; digestion; PCR; primer; se. Human secreted salivary protein zsig63 PCR primer ZC7321.

Homo sapiens.

03-AUG-2001; 2001US-00922480 JS2002081701-A1 27-JUN-2002. 

17-MAR-1999; 99US-0124820P. 17-MAR-2000; 2000US-00527345.

(ADLE/) ADLER D A. (SHEP/) SHEPPARD P O.

Adler DA, Sheppard PO;

WPI; 2002-635468/68.

Novel secreted salivary protein, zsig63 and polynucleotide encoding it useful for treating microbial infections, inflammatory conditions, dental caries and lung infections associated with cystic fibrosis.

Example 1; Page 29; 33pp; English.

The present invention relates to a new secreted salivary protein, zsig63.

The invention is useful for detecting in a test sample, the presence of
an antagonist or agonist of zsig63 protein activity. The invention is
also useful as an immunogen for producing an antibody to zsig63

CC polypeptide. zsig63-cytokine fusion proteins or antibody-cytokine fusion

protein are useful for enhancing in vivo killing of target tissues.

CC periodic composition comprising purified zsig63 polypeptide are

useful in the treatment of conditions associated with pathological

microbes, including bacterial, fungal and viral infections. High

expression of ssig63 in salivary gland suggests that anti-microbial

CC sapplications are useful for treatment of dental caries (tooth decay),

periodontal disease, thrush and gastrointestinal disease. Other

conjugations can be used in uninary tract infections, vaginal infections,

prevention of infection in skin and other epithelial wounds. The

polypeptides can be used in uninary tract infections, vaginal infections,

prevention of infection in skin and other epithelial wounds. The

polypeptides can be used in uninary tract infections, vaginal infections,

prevention of infection in skin and other epithelial wounds. The

polypeptides can be used in uninary tract infections for such pro
inflammatory activity include the treatment of chronic tissue damage,

when pro-inflammatory activity is desired. Applications for such pro
inflammatory activity include the treatment of chronic tissue damage,

collypeptides may be useful as anti-inflammatory agents. The invention is

such as AIDS (acquired immunodeficiency syndroment immune system,

such as AIDS (acquired immunodeficiency syndroment immune system,

such as AIDS (acquired immunodeficiency syndroment in intervention is

that have undergone chemocherapy, radiation treatment. The invention is

also useful for the treatment of lung infections associated with cystic

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fibrosis and its agonists or antagonists are useful for aiding digestion. The present nucleic acid sequence represents a PCR primer that was used in the methods of the invention for identification of zsig63
                                                                                                                                         Gaps
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                                                                                                       Score 25.2; DB 1; Length 26;
                                                                                                                                         0; Indels
                                                                  Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                         3.6e+02;
                                                                                                                                         1; Mismatches
                                                                                                                         Pred. No.
                                                                                                                       96.28;
                                                                                                                     1 Similarity 96.2 25; Conservative
                                                                                                       Query Match
                                                                                                                         Local
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 859938
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2708 TAAAAAAAAAAAAAAAAAAAA 2733 BAAAAAAAAAAAAAAAAAAAAA 1 26 ઠે g

BP

AAD45054 standard; DNA; 26

AAD45054/c

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Human; secreted salivary protein; zsig63 protein; host defense protein; immune modulating factor; antipathogenic; cell-cell signalling molecule; growth factor; oytokine; growth factor hormone activity; dental carries; infection; tooth decay; periodontal disease; gastrointestinal disease; thrush; urinary tract infection; vaginal infection; diabetes; obesity; anti-inflammatory; chronic tissue damage; lung dysfunction; restenosis; gene therapy; salivary gland dysfunction; prostate gland dysfunction; forensic DNA profiling; chondrosarcoma; atherosclerosis; primer; ss.
                                                 ZC7231 primer used in the identification of human zsig63 DNA.
                             (first entry)
                             27-DEC-2002
         AAD45054;
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03-AUG-2001; 2001US-00923236. US2002090677-A1. Homo sapiens 11-JUL-2002.

17-MAR-1999; 99US-0124820P. 17-MAR-2000; 2000US-00527345. (ADLE/) ADLER D A. (SHEP/) SHEPPARD P O.

Sheppard PO; Adler DA,

WPI; 2002-642378/69.

Novel secreted salivary polypeptide, zsig63, useful as antimicrobial agent for treating microbial infection, dental carries, periodontal disease, thrush gastrointestinal disease, and for aiding digestion.

Example 1; Page 29; 33pp; English.

The invention relates to human secreted salivary polypeptide designated as zaig63 and nucleic acid molecules encoding such polypeptides. zaig63 can be used in detecting agonists and antagonists of its activity, and is also useful as a host defense polypeptide, immune modulating factor, antipathogenic polypeptide, cell-cell signalling molecule, growth factor, cytokine, or as secreted extracellular matrix associated protesins with growth factor hormone activity. It is useful for treating conditions associated with pathological microbes, including bacterial, fungal and cyral infections, for treating dental carries (tooth decay), periodontal clisease, thrush and gastrointestinal disease, for treating urinary tract infection, vaginal infection and for preventing infection in skin and controllors and protect against pathogenic colonisation and invasion, for treating chronic tissue damage erg. damage in extremities associated with thabetes and useful as anti-inflammatory agents. It is useful as marker

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  periodontal disease; thrush; gastrointestinal disease; epithelial wound; periodontal disease; thrush; gastrointestinal disease; epithelial wound; urinary tract infection; vaginal infection; skin infection; primer; pro-inflammatory; chronic tissue damage; vascular system; diabetes; AIDS; lung infection; cystic fibrosis; lung dysfunction; digestive; salivary gland carcinoma; Pneumocystis carinii infection; emphysema; chronic bronchitis; prostate dysfunction; prostate adenocarcinoma; cell culture media; gene therapy; human chromosome 4q12-4q13; dentinogenesis imperfecta; dentin dysplasia type II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated zsig63 polypeptide comprising at least 90% identity to an amino acid sequence which comprises domain 1 of zsig63, domain 2, domain 3, maturez sig63 and full length zsig3. Also included are the polymucleotide encoding zsig63, a zsig63 expression vector, a cultured cell comprising the vector and expressing the protein,
of lung dysfunction, salivary gland dysfunction, or dysfunction of prostate gland. It is also therapeutically useful for aiding digestion. Polynucleotides of the invention are used in gene therapy for increasing or inhibiting zsig63 activity, for detecting abnormalities on human chromosome 4 associated with disease or other human traits and as diagnostics in forensic DNA profiling. Sequences of the invention are useful for stimulating proliferation or differentiation of cardiac myocytes, for proliferation or differentiation of addiacting chandrosarcomas, atherosclerosis, restenosis and obesity. The present sequence is a primer used in the identification of human zsig63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated zsig63 polypeptide, member of the adhesin family, useful for treating dental carries, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections,
                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                             0.9%; Score 25.2; DB 1; Length 26; 96.2%; Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                  Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human zaig63 PCR/sequencing primer ZC7231
                                                                                                                                                                                                                                                                                                                                                                            2708 TAAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                     ВААААААААААААААААААААА 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABX93598 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-MAR-1999; 99US-0124820P.
17-MAR-2000; 2000US-00527345.
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                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ADLE/) ADLER D A. (SHEP/) SHEPPARD P O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-328428/31.
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                                                                                                                                                                                                                                                                                                                                     25;
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                                                                                                                                                                                                                                                                                               Query Match
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a DNA encoding a fusion protein (comprising amino acids 1-15, 16-37, 38-16, 127-219 or 16-219 of zsig63 and an additional protein), using a zsig63 reporter gene construct to identify zsig63 reporter gene construct to identify zsig63 and producing an anti-zsig63 antibody using zsig63 immunogenic peptides, Zsig63 is useful for detecting in a test sample, the presence of antagonist of useful for detecting in a test sample, the presence of antagonist of catig63 protein activity. Zsig63 has antimicrobial activity and since exhibits high expression in salivary thrush, and gastrointestinal disease, urinary tract infections, vasin infections and other epithelial wounds. The polypeptides can be used for treating disease, urinary tract infections, vasin infections and other epithelial wounds. The polypeptides can be used to establish normal microflora and protect against pathogenic colonization and civity for treating chronic, tissue damage particularly in areas having limited or tamaged vascular system, e.g. in diabetes, and for treating limited or democherapy, radiation treatment, for treating ung infections e.g. in chemotherapy, radiation freatment, for treating ung infections e.g. in cystic fibrosis. Detection of zsig63 polypeptide at relatively high cystic fibrosis. Detection of zsig63 polypeptide at relatively high marker of lung dysfunction in Saig63 polypeptide at relatively high conditions associated with salivary gland or lung dysfunction and soconditions and and may be used to replace serum that is commonly used in culture media and may be used to replace serum that is commonly used in culture media and may be used to replace serum that is commonly used in charged and may be used to replace serum that is commonly used continue and and may be used to replace with the tasing imperfects, and dencin dysplasia type II). Zsig63 is an adheshin family member. The continue appears the engage of isolate and sequence is a primer used to isolate and sequence nucleic acdis encoding human zsig63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid manipulation, mRNA profiling, polymerase chain reaction,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 25.2; DB 1;
96.2%; Pred. No. 3.6e+02;
tive 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Metsis A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of a second back primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    electrophoresis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
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Best Local Similarity
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Indels

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1; Mismatches

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25; Conservative
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                                                                                                                                                                                                                                                                                                                      primer; 88.
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                                                                                                                               ADY96656;
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                                                                                  RESULT 218
   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for the prophylaxis infections, comprises
profiling, comprises amplification by nested polymerase chain
                                                                   product DNA molecules comprising amplification by a nested PCR method. The method is useful in profiling mRNA transcribed in a system under investigation. The oligonuclectides are used as size standards in electrophoresis, and as intercond controls allowing for calculation of relative amounts of material present. The present sequence represents specific example of a PCR primer used in the method of the invention
                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                            Bovine Viral Diarrhea Virus; BVDV; infection; vaccine; prophylaxis;
                                                         invention relates to producing a population of double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                       Bovine viral diarrhea virus gene 5' end amplifying PCR primer.
                                                                                                                                                                         DB 1; Length 26;
                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New DNA molecule useful for manufacturing a vaccine and treatment of Bovine Viral Diarrhea Virus (BVDV) a sequence complementary to a BVDV RNA.
                                                                                                                                                  Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                        Score 25.2; DB 1;
Pred. No. 3.6e+02;
1; Mismatches 0
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                                 Claim 6; Page 85; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; PCR; primer; ss.
                                                                                                                                                                         0.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-2002; 2002WO-EP009925
                                                                                                                                                                                                                                                                                             692/c
AAD55692 standard; DNA; 26
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                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-333043/31
                                                                                                                                                                                    Local Similarity
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07-AUG-2003
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                                                                                                                                                                                                                                                                                                                              AAD55692;
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Score 25.2; DB 1; Length 26; Pred. No. 3.6e+02;

0.9%; Score 25.2; 96.2%; Pred. No. 3.

Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to zsig63, a novel secreted salivary protein and its encoding polynucleotide. Zsig63 is a member of the adhesin family. The invention is useful for diagnosing and treating microbial infections such as dental carries, periodontal disease, thrush, gastrointestinal disease, uninary tract infection, vaginal infection, skin infection or epithelial wounds and lung disfunctions such as AIDS, lung infections and cancer. The invention is also useful in gene therapy. The present sequence is human Zsig63 cDNA cloning and sequencing primer. This sequence to obtain the full-length Zsig63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   wer polynucleotide (I) encoding a zaig63 polypeptide, useful for diagnosing and treating microbial infections, e.g. dental carries, thrush, gastrointestinal disease, skin infection, or epithelial wounds, AIDS, lung infections, or cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human Zsig63 cDNA cloning and sequencing primer, ZC7231.
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2708 TAAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                      26 BAAAAAAAAAAAAAAAAAAA 1
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17-MAR-2000; 2000US-00527345.
03-AUG-2001; 2001US-00923236.
                                                                                                                                                                                                                                                             ADY96656 standard; DNA; 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sheppard PO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ò
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Transgenic mammal or its embryo useful as model for human disease, has heterologous nucleotide sequence coding for stratum corneum chymotryptic enzyme operably linked to promoter that drives its expression in skin.
                                                                                            SCCE; murine; stratum corneum chymotryptic enzyme; kallikrein 7; serine protease; transgenic mammal; skin; skin disease; skin cancer; hyperkeratosis; acanthosis; epidermal inflammation; dermal inflammation; pruritus; atopic dermatitis; eczema; acne; itch; KLK7; ss.
                                                                           Murine SCCE 5'-RACE oligonucleotide SEQ ID 42.
                                                                                                                                                                                                                                                                                                                                                                Example 6; Page 36; 74pp; English.
                   ABQ76254 standard; DNA; 27 BP
                                                                                                                                                                                                       08-FEB-2002; 2002WO-IB001300
                                                                                                                                                                                                                          09-FEB-2001; 2001CA-02332655
09-FEB-2001; 2001DK-00000218
                                                         (first entry)
                                                                                                                                                                                                                                                                                      Sgelrud T, Hansson L;
                                                                                                                                                                                                                                                                                                       WPI; 2002-643380/69
                                                                                                                                                                                                                                                       (EGEL/) EGELRUD T. (HANS/) HANSSON L.
                                                                                                                                                                  WO200262135-A2.
                                                                                                                                                Mus musculus
                                                         08-NOV-2002
                                                                                                                                                                                     15-AUG-2002.
                                      ABQ76254;
RESULT 219
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combryo having integrated within its genome, a heterologous nucleotide embryo having integrated within its genome, a heterologous nucleotide embryo having integrated within its genome, a heterologous nucleotide sequence comprising at least a significant part of a nucleotide sequence coding for a stratum corneum chymotryptic enzyme (SCCE) or its variant, coding for a product of the invention is useful as a model or the study of disease with the aim of improving treatment, to relieve for the study of disease with the aim of improving treatment, to relieve commentic or a pharmaceutical formulation, and for the development of a disgnostic method. It can also be used as a model for a skin cancer. The invention is also useful for screening or identifying a compound or composition effective for the prevention or treatment of an abnormal or unwanted phenotype, and for screening or identifying a compound or composition effective for the prevention or treatment of an abnormal or unwanted phenotype, and for screening or identifying a compound or composition effective for the prevention or treatment of an abnormal atopic dermatitis, eczema, accessisting of epidermal hyperkeratosis. The mammal of the invention is also useful as a compounds and compositions for relieve of various skin diseases where itch is a component. This sequence represents a 5' RACE cDNA synthesis primer used in a method of detecting homologues to human caracturing and the homotryptic enzyme, SCCE, gene. SCCE is a serine stratum corneum chymotryptic enzyme, SCCE, gene. SCCE is a serine protease synonymous with human kallikrein 7 (KLK7) and is used in the development of the transgenic mammals described in the invention

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Sequence 27 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 2 Other;
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Query Match 0.9%; Score 25.2; DB 1; Length 27; Best Local Similarity 96.2%; Pred. No. 3.7e+02; Matches 25; Conservative 1; Mismatches 0; Indels
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The invention relates to specific expressive sequential labels of tubulin of tea tree and their biochips. The invention can be used in evaluation of crop seed sources, early prediction of hybrid vigor, research on plant resistance, determination of plant SNP, inspection of transgenic crop security, and soreen of herbicides and agro-chemicals, etc. The present sequence represents a tea tree tubulin oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Tubulin differential expression sequence label of tea tree and biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide biotin-T25 for novel nucleic acid immobilisation method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immobilisation; solid support; salt; cationic detergent; capture probe;
hybridisation; primer; template-dependent extension; target organism;
sequencing; genetic polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 27 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 . Match 0.9%; Score 25.2; DB 1;
Local Similarity 96.2%; Pred. No. 3.7e+02;
les 25; Conservative 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2708 TAAAAAAAAAAAAAAAAAAAAAA 2733
2708 TAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tea tree tubulin oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                  (TEAC-) TEA INST CHINESE AGRIC ACAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Page 19; 25pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ95960 standard; DNA; 25 BP.
                                                                                                                                                                                                                                                                                                              18-DEC-2003; 2003CN-01109578.
                                                                                                                                                                                                                                                                                                                                      18-DEC-2003; 2003CN-01109578.
                                                                                           AEA37617 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                             Xu Y, Zhao L;
                                                                                                                                                                                                                                Melaleuca alternifolia
                                                                                                                                                                                                    ss; tubulin; biochip.
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2005-197097/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-FEB-1996
                                                                                                                                                 28-JUL-2005
                                                                                                                                                                                                                                                          CN1552861-A.
                                                                                                                                                                                                                                                                                    08-DEC-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ95960;
                                                                                                                       AEA37617;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                             Chen L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                    RESULT 220
                                                                                  AEA3761
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13-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-2000
                                                                                                                                                                                                                                                                                                                                                 Synthetic.
           Sakaki Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wages JM,
                                                                                                                                                                                                                                                                   AAA39306;
                                                                                                                                                    Query Match
Best Local S
                                                                                                                                                                                                                               RESULT 223
AAA39306/c
                                                                                                                                                                      Matches
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                                                                                                                                                                                                                            Oligonucleotides AAQ95959-82 are examples of oligonucleotides used in a novel method of immobilising oligonucleotides to a solid support by incubating in the presence of a salt or cationic detergent e.g. NaCl (50-250 mM, pH 6.0-8.0) or 1-ethyl-3-(3'-dimethyl amino propyl)-1,3 carbodiimide hydrochloride (ECD). The oligonucleotides can be capture probes for detection of specific nucleic acids by hybridisation or can be primers for template-dependent extension from the immobilised primers on nucleic acid from a target organism. The method can be used in hybridisation assays, sequencing and analysis of genetic polymorphism
                                                                                                                                                                         Immobilising synthetic nucleic acid on solid support - by incubation in presence of salt or cationic detergent, for use in hybridisation assays, sequencing and analysis of polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein 1; Napl; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer for human Nck associated protein 1 coding sequence.
                                                                                                                                                                                                                                                                                                                                     0.9%; Score 25; DB 1; Length 25; 100.0%; Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                   Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                         100.0%; Preu. ...
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                 /*tag= a
/note= "biotinylated"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 18; 61pp; English.
                                                                                             93US-00162397
94US-00341148
                                                                            94WO-US014096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98WO-JP005646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KYOW ) KYOWA HAKKO KOGYO KK.
(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97JP-00363183
                                                                                                                     (MOLE-) MOLECULAR TOOL INC
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX84259 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy; PCR primer; ss.
                                                                                                                                        Nikiforov T, Knapp MR;
                                                                                                                                                          WPI; 1995-224282/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        associated
         misc_feature
                                                                            06-DEC-1994;
                                          WO9515970-A1
                                                                                              06-DEC-1993;
                                                                                                      16-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-DEC-1998;
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                                                          15-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                        25
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                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 222
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The present invention describes a rapid pairing reagent (I) for the isolation or detection of a polymucleotide (PN) analyte molecule having a selected target base sequence, in a sample containing the analyte molecule and non-target polymucleotide, comprising a capture component (A) and a target-specific probe (B) linked to a solid substrate. The isolated sequences are useful for enzymatic amplification. (I) is capable of rapidly binding nucleic acids in the sample and placing them in close proximity to target probes on the reagent, thus enabling binding under low stringency. Combination of rapid capture and concentration of polymucleotides with selective targeting of analyte molecules, greatly enhances the isolation process. Non-ionic morpholino oligomers used as probes are not extended by polymerases and therefore do not interfere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reagent for rapidly detecting or isolating target nucleic acid sequences in polynucleotide-containing sample, comprises capture component and target-specific probe linked to solid substrate.
                                                                                                                                                                                                                                                              This sequence represents a PCR primer used to isolate DNA encoding the human Nck associated protein 1 (Napl) of the invention. Napl inhibits apoptosis. The protein can be used in the investigation, diagnosis and treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                      Protein inhibiting apoptosis, useful in the diagnosis and treatment of Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rapid detection; probe; target nucleic acid; enzymatic amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25; DB 1; Length 25;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rapid capture probe designated Neu-probe SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 1 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.9%; Sco...
100.0%; Pred. No. ...
0; Mismatches
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                                                                                                                                                                                           Disclosure; Page 76; 90pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Weller DD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Col 17; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-00969813.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA39306 standard; RNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    isolation; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Summerton JE,
WPI; 1999-395181/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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The present sequence represents a capture probe specific for a c-myc fusion target. It is used in the method of the invention. The specification describes the use of non-nucleoside spacers to immobilise an array of capture probes on a solid support. The solid support carries an array of capture probes, each consisting of non-nucleoside spacers on a notigonucleotide to which a nucleic acid-protein fusion or a ribons an oligonucleotide to which a nucleic acid-protein fusion or a ribons with the support surface, ensuring efficient hybridisation between capture probes and bound nucleic acid/protein this minimaling denaturation of the protein which may then companies, while minimaling denaturation of the protein which may then compared in interactions between proteins and compounds (e.g. other proteins, ligands or nucleic acids), particularly to identify potential therapeutic agents, ensyme substrates or unknown proteins that interact with drugs, but also for diagnosis (detecting disease-associated proteins) and for quantifying target molecules in a sample
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with amplification of target molecule. AAA39316 to AAA39316 represent oligonucleotides used in the exemplification of the present invention
                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       c-myc fusion; non-nucleoside spacer; capture probe;
nucleic acid-protein fusion; ribosome display particle; ss.
                                                                                                                                                 DB 1; Le...
0. 3.7e+02; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capture probe CP125 specific for c-myc fusion targets.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                    Sequence 25 BP; 0 A; 0 C; 0 G; 0 T; 25 U; 0 Other;
                                                                                                                                                                           Query Match 0.9%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 3.7 Matches 25; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ30267 standard; DNA; 25 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-013048/01.
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The invention relates to a method of preparing a controlled pore glass-polytetrafluoroethylene (PTE) resin chromatography membrane, comprising combining controlled pore glass and an aqueous dispersion of PTFE to form a paste-like mass, heating the paste-like mass at 50-70 plus oC, calendering to form a foldable sheet, and sintering the sheet to produce a rigid, porous sheet. The method prepares a controlled pore glass-PTFE resin chromatography membrane for use in various biotechnical procedures. The membrane is useful in place of controlled pore glass as a support for the synthesis, isolation, and purification of nucleic acids and for the isolation and purification of proteins. The method produces a membrane that may be used in lieu of controlled pore glass. The present sequence trapresents an oligonucleotide prepared on the membrane in an example which demonstrates the method of the invention
                                                                                                                                                                                                     Glass-synthetic resin membrane; pore glass-polytetrafluoroethylene resin;
chromatography membrane; PTFE; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Preparation of controlled pore glass-polytetrafluoroethylene resin chromatography membrane by heating, calendering and sintering mixture of controlled pore glass and aqueous dispersion of polytetrafluoroethylene.
                                                                                                                                                                        Example oligonucleotide #2 prepared on glass-synthetic resin membrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide of the invention SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ААААААААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 12; Col 8; 6pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                        99US-00305219
                                                                                                                                                                                                                                                                                                                                                                                        96US-00604440.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC54009 standard; DNA; 25
                                                                            ABK49986 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-534961/59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen R;
                                                                                                                                                                                                                                                                                                                                                                                                                      (CPGC-) CPG INC
                                                                                                                                              15-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                        04-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                          21-FEB-1996;
                                                                                                                                                                                                                                                                                            US6261497-B1
                                                                                                                                                                                                                                                                                                                          17-JUL-2001.
                                                                                                                                                                                                                                                            Synthetic.
                                                                                                              ABK49986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wong YN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
   25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                  RESULT 22:
ABK49986/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC54009,
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25; Conservative

Best\_Local\_Similarity Matches 25; Conserv

Query Match

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Gaps

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0; Indels

0.9%; Score 25; DB 1; Length 25; 100.0%; Pred. No. 3.7e+02;

Gaps

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The invention relates to a method and device for the manufacture of a probe array. The method involves using an inkjet system to discharge a probe solution through a solution discharging head, so as to form a number of probes on a solid matrix. Mists of the probe solution generated during probe solution discharge are caught by an electrostatic adeorption mechanism. The method and device are suitable for manufacturing probe arrays for analysing DNA sequences, and for the simultaneous detection of multiple genes. The method and device of the invention prevent the scattering of probe positions and the mixing of different probe solutions. The present sequence is related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Probe array; microarray; DNA array; DNA chip; manufacture; inkjet system; electrostatic adsorption mechanism; DNA analysis; simultaneous gene detection; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Manufacturing of probe carrier for carrying probes for base sequence analysis of genetic deoxyribonucleic acid and simultaneous multiple item diagnosis of gene by ink jet process while removing mist of probe
solution, and removes liquid droplets adhering to discharge nozzle. The present sequence is used in the exemplification of the invention.
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                                                                                                       0.9%; Score 25; DB 1; Length 25;
100.0%; Pred. No. 3.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.9%; Score 25; DB 1; Length 25; 100.0%; Pred. No. 3.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe #4, immobilised on probe array using novel method.
                                                              Sequence 25 BP; 25 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                          2709 AAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                                                                                 AAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                            ADF39737 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                    25; Conservative
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                                                                                                                             Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2003014773-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                         ADF39737;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  solution.
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                                                                                                         Query Match
Best Local
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ADF39736
                                                                                                                                                                                                                                                                                                    RESULT 228
                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                              Probe carrier manufacturing method for inkjet system, involves scanning liquid discharge head in direction orthogonal to scanning direction, at angle satisfying predetermined relation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel probe carrier and the method for manufacturing the carrier. The invention enables stable discharge of solution, and removes liquid droplets adhering to discharge nozzle. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Probe carrier manufacturing method for inkjet system, involves scanning liquid discharge head in direction orthogonal to scanning direction, at angle satisfying predetermined relation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel probe carrier and the method for manufacturing the carrier. The invention enables stable discharge of
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.9%; Score 25; DB 1; Length 25; 00.0%; Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide of the invention SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.0%; Pred. No. 3.7
Matches 25; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 4; 17pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC54008 standard; DNA; 25 BP
               ss; probe carrier; discharge.
                                                                                                                                                                                   28-MAR-2002; 2002JP-00093023.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-MAR-2001; 2001JP-00094400.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; probe carrier; discharge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAR-2002; 2002JP-00093023
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                                                                                                                                                                                                                                                                                                                   WPI; 2003-535999/51.
                                                                                                                                                                                                                                                                           (CANO ) CANON KK.
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                                                                                                JP2003035711-A.
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                                                                                                                                           07-FEB-2003
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                                                     Synthetic.
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Gaps

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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA sample containing at least one segment of (1); and determining the length of the resulting amplicon(s). Also described are:

The method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a disease related gene, that are associated with a predisposition to disease and for prediagnosis of such diseases, especially prion diseases but also cype genes that encode milk proteins, classases and for prediagnosis of such diseases, aspecially prion diseases because or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a prion protein polymorphic microsatellite marker
                                                                                                                                                                                                                                                                                                      Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fluorophore; detection; antibody; antigen; avidin; hormone; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fluorophore-labeled biological detection oligonucleotide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.9%; Score 25; DB 1; Length 25;
100.0%; Pred. No. 3.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MONS ) MONSANTO TECHNOLOGY LLC.
                                                                                                                                                                                                                                                                                                                                                                                                           Claim 9; Page 50; 64pp; German.
                                                                                                                                                                                                                          Han
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-MAY-2001; 2001US-00859736.
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                                                                                    09-AUG-2002; 2002DE-01036711.
                                                                                                                                 09-AUG-2002; 2002DE-01036711.
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                                                                                                                                                                           (UYHO-) UNIV HOHENHEIM
                                                                                                                                                                                                                        Preuss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                  WPI; 2004-215730/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      consensus sequence.
                                                                                                                                                                                                                        Geldermann H,
DE10236711-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6838244-B1.
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                                        26-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method and device for the manufacture of a probe array. The method involves using an inkjet system to discharge a probe solution through a solution discharging head, so as to form a number of probes on a solid matrix. Mists of the probe solution generated during probe solution discharge are caught by an electrostatic adsorption mechanism. The method and device are suitable for manufacturing probe arrays for analysing DNA sequences, and for the simultaneous detection of multiple genes. The method and device of the invention probe scattering of probe positions and the mixing of different probe solutions. The present sequence is related to the invention.
                                                                                                                                                                              DNA chip, manufacture, inkjet system;
DNA analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Manufacturing of probe carrier for carrying probes for base sequence analysis of genetic deoxyribonucleic acid and simultaneous multiple item diagnosis of gene by ink jet process while removing mist of probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion protein polymorphic microsatellite marker consensus sequence #23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; malk protein; hormone; transcription factor; pT7-blue-vector; sheep;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                   Target DNA sequence #3, capable of hybridising to probe #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seguence 25 BP; 25 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 25; DB 1; Le
100.0%; Pred. No. 3.7e+02;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; SEQ ID NO 3; 15pp; Japanese
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                                                                                                                                                                              DNA array;
mechanism;
                                                                                                                                                                              Probe array; microarray; DNA arr
electrostatic adsorption mechani
simultaneous gene detection; ss.
    ADF39736 standard; DNA; 25 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-496695/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                microsatellite; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CANO ) CANON KK.
                                                                                                                                                                                                                                                                                                                      JP2003014773-A.
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                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADO81145;
                                            ADF39736;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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Best Loca Matches

ð 유 RESULT 230 AD081145,

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Gaps

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Zhou JS;

Li WR,

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The invention relates to an oligonucleotide molecule (ON) labeled with several fluorophores of one or more types embedded in its backbone, where one or more of fluorophores is not located at either the 3' or 5' terminus of ON. ON is useful for sequencing nucleic molecules. ON is useful for detecting biological molecules e.g., antibody, antigen, avidin, protein, peptide, bacteria, virus, blood cell or hormone. ON is capable of providing strong fluorescence signals at different wavelengths. This sequence corresponds to an example of an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an oligonucleotide molecule (ON) labeled with several fluorophores of one or more types embedded in its backbone, where one or more of the fluorophores is not located at either the 3' or 5' terminus of ON. ON is useful for sequencing nucleic molecules. ON is useful for detecting biological molecules e.g., antibody, antigen, avidin, protein, peptide, bacteria, virus, blood cell or hormone. ON is capable of providing strong fluorescence signals at different
                                                                    oligonucleotide molecule labeled with several fluorophores, useful detecting biological molecules e.g., antibody, antigen, avidin or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel oligonucleotide molecule labeled with several fluorophores, useful for detecting biological molecules e.g., antibody, antigen, avidin or
                                                                                                                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fluorophore; detection; antibody; antigen; avidin; hormone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fluorophore-labeled biological detection oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        wavelengths. This sequence corresponds to an example of an
                                                                                                                                                                                                                                                                                                                                                                                Score 25; DB 1; Length 25; Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                    Example 1; SEQ ID NO 2; 18pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Scur
100.0%; Pred
0; M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MONS ) MONSANTO TECHNOLOGY LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADV86468 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAY-2000; 2000US-0205452P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-MAY-2001; 2001US-00859736.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-MAR-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  25; Conservative
                                                                    Novel oligonucleotide
                                  WPI; 2005-063191/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2005-063191/07.
                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6838244-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-JAN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                      protein.
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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "Optionally conjugated via 5' phosphate to (CH2)6
linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Manufacture of probe carrier for analyzing gene deoxyribonucleic acid sequence, involves forming labeled indexes on carrier at specific postitions, and applying solutions respectively containing probes to respective specific positions.
                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method for the manufacture of a DNA probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'bound_moiety= "Bases 25-1 of target SEQ ID NO:3"
                                                                                               ö
                                                                 Length 25;
                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                 DNA microarray; biochip; immobilization; probe; ss.
                                T; 0 U; 0 Other;
                                                           0.9%; Score 25; DB 1; I
100.0%; Pred. No. 3.7e+02;
ive 0; Mismatches 0;
                                                                                                                              2709 AAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 4; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                DNA hybridization probe, SEQ ID NO:4.
                                                                                                                                                             25 AAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Socation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kameyama M;
                               Sequence 25 BP; 0 A; 0 C; 0 G; 25
oligonucleotide of the invention.
                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-OCT-2004; 2004US-00958348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-APR-2001; 2001JP-00133697
27-APR-2001; 2001JP-00133698
29-APR-2002; 2002US-00133675
                                                                                                                                                                                                                                                 AEB26392 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                            1 Similarity 100.
.25; Conservative
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_binding
                                                                                                                                                                                                                                                                                                                 22-SEP-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                  AEB26392;
                                                              Query Match
Best Local
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                                                                                               Matches
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method can provide an array substrate that prevents probe solutions being applied at adjacent sites from being mixed with each other and allows the probe solutions to sufficiently spread in the respective regions to prevent blank areas from being produced, thereby permitting the efficient production of reliable probe arrays. Sequences AEB26389-AEB26393 represent oligonicalcotides used to illustrate the invention. The present sequence represents a probe which is complementary to the target shown as SEQ ID NO:3 (AEB26391) which was immobilized on a substrate according to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Manufacture of probe carrier for analyzing gene deoxyribonucleic acid sequence, involves forming labeled indexes on carrier at specific positions, and applying solutions respectively containing probes to
                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               bound_moiety= "Bases 25-1 of probe SEQ ID NO:4"
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/note= "Optionally conjugated to fluoresent dye
tetramethylrhodamine"
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                                                                                                                                                                                          Length 25;
                                                                                                                                                                                                                       0; Indels
                                                                                                                                                           Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                        DB 1; Le
3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fluorescently-labeled DNA target, SEQ ID NO:3.
                                                                                                                                                                               0.9%; Scor.
100.0%; Pred. No. ...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ONA microarray; biochip; immobilization; ss
                                                                                                                                                                                                                                                      2709 AAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; SEQ ID NO 3; 24pp; English.
                                                                                                                                                                                                                                                                                  25 AAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kameyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      respective specific positions.
                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-APR-2001; 2001JP-00133697.
27-APR-2001; 2001JP-00133698.
29-APR-2002; 2002US-00133675.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-OCT-2004; 2004US-00958348
                                                                                                                                                                                                                                                                                                                                                                 AEB26391 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 '*tag≔ a
                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JS2005158738-A1
                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_binding
                                                                                                             SEQ ID NO:3
                                                                                                                                                                                                                                                                                                                                                                                                                             22-SEP-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                AEB26391;
                                                                                                                                                                                                                                                                                                                                   RESULT 234
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cc further comprises forming a dividing wall (especially by photolithography cc methods) for partitioning the specific positions on the substrate, irradiating the substrate with plasma in a gas atmosphere containing fluorine, and removing ingredients of the gas adhering to the specific positions. The invention also relates to a method of identifying the contining the invention also relates to a method of identifying the contining the position of a target substance bonded to the probe on such a substrate by cutilizing the indexes. The probe array is used for analyzing gene capplied at adjacent sites from being multiple genetic diagnoses. The inventive completed at adjacent sites from being mixed with each other and allows the probe solutions to sufficiently spread in the respective regions to probe colutions to sufficiently spread in the respective regions to production of reliable probe arrays. Sequences AEB26393 AEB26393 Crepresent oligonucleotides used to illustrate the invention. The present sequence represents a tetramethylrhodamine-labeled model target sequence which is complementary to the probe shown as SEQ ID NO:4 (AEB26392).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or detecting,
the breast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BS124; breast; cancer; detection; diagnosis; prevention; treatment; EST;
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4, Colpitts TL, Friedman PN, Gordon J;
Klass MR, Kratochvil JD, Russell JC;
Yu H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of an oligonucleotide used in the isolation of BS124-specific EST clone. It is useful for detecting, diagnosing, staging, preventing or treating, or determining predisposition to diseases or conditions of the breast, such as breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated BS124 polynucleotides and polypeptides - used for diagnosing, preventing or treating diseases or conditions of the
                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                         Score 25; DB 1; Length 25; Pred. No. 3.7e+02;
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                                                                                                                                                                                                                                                                                                                                     Sequence 25 BP; 25 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 26 BP; 0 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human BS124 specific EST clone oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Prec. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Billing-Medel PA, Cohen M,
Granados EN, Hodges SC, Kl
Scheffel CP, Stroupe SD, Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX07466 standard; cDNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              such as breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-105623/09
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Granados EN,
Scheffel CP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX07466;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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AAX07466/
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RESULT 236

AAX78723,

Matches

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This sequence represents a molecular scaffold which may be used in a composition of matter comprising multiple fluorophores. The fluorophores are bound to the molecular scaffold at separate predetermined positions, to permit fluorescence energy transfer between two fluorophores. The fluorophores are characterized by maximum emission wavelength of one being greater than the minimum excitation wavelength of the other. The composition is useful for determining whether a preselected nucleotide residue is present at a predetermining whether a preselected nucleotide residue is present at a predetermined position within a nucleic acid. It is also useful in multicomponent analysis including multiplex biological analysis, and identifying multiple single nucleotide polymorphisms. The presence of a number of given nucleotide residues is determined simultaneously by the composition of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Composition of matter useful for multi-component analyses, comprises multiple fluorophores bound to molecular scaffold at preset positions to permit fluorescence energy transfer between two fluorophores.
                                                                                                 Molecular scaffold; fluorophore; fluorescence; energy transfer; emission wavelength; excitation wavelength; multiple; single nucleotide; polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; B-cell proliferation; anti-tumour response; immune system; immunostimulant; cytostatic; human; sequencing primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.9%; Score 25; DB 1; Length 26;
100.0%; Pred. No. 3.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human zalphall Ligand sequencing primer ZC7764b.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Pred. No. 5...
Marches 25; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Russo JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                             11-SEP-2000; 2000US-00658077. 31-JUL-2001; 2001US-0309156P.
                                                                                                                                                                                                                                                                                                                                    11-SEP-2001; 2001WO-US028967.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS20672 standard; DNA; 26
         (first entry)
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                                                    Scaffold oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ju J, Li Z, Tong A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-575158/61.
                                                                                                                                                                                                                                          WO200222883-A1.
           24-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6307024-B1
                                                                                                                                                                                                                                                                                       21-MAR-2002
                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS20672;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel contiguous and partially overlapping cDNA sequences and their encoded polypeptides, designated PA153, transcribed from human pancreatic tissue and which have cytostatic activity. The PA153 polymucleotides, proteins and antibodies are all useful in methods of detection. Detection of PA153 polymucleotide, antigens or anti-PA153 antibodies in a sample is indicative of pancreatic disease. PA153 antibodies (antagonists) can also be used in vivo for therapeutic use, e.g. treatment of pancreatic disease, tumours or metastases. Antisense PA153 polymucleotides can be used in gene therapy of pancreatic diseases. AAX78712-X78725 represent primers used in the method of the invention
                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Billing-Medel PA, Cohen M, Colpitts TL, Friedman PN, Gordon J;
Granados EN, Hodges SC, Klass MR, Kratochvil JD, Roberts-Rapp L;
Russell JC, Stroupe SD;
                                                                                                                                                                                                                                                                                                                                                                                                              Pancreatic disease; PA153; human; cytostatic; detection; antigen; anti-PA153; antagonist; therapy; treatment; tumour; metastasis; gene therapy; EST; expressed sequence tag; primer; ss.
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                            Indels
                                                                                                                                                                                                                                                                                                                                                                          Human pancreatic PA153 EST-specific clone primer 12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26 BP; 0 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
       3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PA153 cDNA transcribed from pancreatic tissue.
    1 Similarity 100.0%; Pred. No. 3.7
25; Conservative 0; Mismatches
                                                                       2709 AAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                25 AAAAAAAAAAAAAAAAAAAAAA
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AAI73048 standard; DNA; 26 BP.
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                                                                                                                                                                                                                                  AAX78723 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                             (first entry)
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9931274-A2
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                                                                                                                                                                                                                                                                                                                             03-SEP-1999
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Gaps

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AAI73048;

AAI73048/c ID AAI73 XX AC AAI73 XX

RESULT 237

8 g 97US-0048810P

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05-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                             25;
                                                                                                                            Russell JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novak JE,
Gross JA,
                                                                                                               Billingel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH44609;
                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                     (FRIE/)
(KLAS/)
(RUSS/)
(STRO/)
                      (BILL/)
                                                                                                                                                                                                                                                                                                          clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                              240
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                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                  The present invention relates to the isolation of a novel cytokine, zalphall Ligand and the polynucleotide encoding it. The invention also gives the sequence for the zalphall receptor and the polynucleotide encoding it. The zalphall Ligand polypeptide stimulates proliferation of attural killer (KK) cells or NK cell progenitors, the activation of natural killer (KK) cells or NK cell progenitors, the activation of NK cells proliferation of F-cells proliferation of B-cells stimulated with anti-IgM antibodies. The zalphall Ligand polypeptide is also useful in preparing antibodies that bind to zalphall Ligand epitopes. The zalphall Ligand polypucleotides can be used as probes or primers to clone regions of a zalphall Ligand gene, and in gene therapy. Zalphall Ligand may also be used to identify inhibitors of its activity, to enhance the generation of anti-tumour responses with or without the influsion of donor lymphocytes, and to activate or stimulate the immune system. The present sequence represents a sequencing primer used to sequence cDNA clones in the isolation of human zalphall Ligand
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                                                                                                                                                                                                  imulating proliferation, activation, differentiation and/or induction inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LS147-specific polynucleotide sequencing related universal primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LS147; cancer; lung cancer; gene therapy; cytostatic; ss; sequencing;
primer; EST clone; expressed sequence tag clone.
                                                                                                                                                                                     zalphall ligand polypeptides and polynucleotides, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.9%; Score 25; DB 1; Length 26; 100.0%; Pred. No. 3.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                            Sprecher CA, Foster DC, Holly Nelson AJ, Dillon SR, Hammond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25 AAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                 Example 7; Col 139; 105pp; English.
                                                        99US-0123547P.
99US-0123904P.
99US-0142013P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВЪ
                                   09-MAR-2000; 2000US-00522217.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.9°
Best Local Similarity 100.
Matches 25; Conservative
                                                                                                      (ZYMO ) ZYMOGENETICS INC
                                                                                                                            Novak JE, Presnell SR,
Gross JA, Johnston JV,
                                                                                                                                                               WPI; 2002-040208/05
                                                                                                                                                                                                                          antigenic response.
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                                                                                                                                                                                               stimulating I
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                                                         09-MAR-1999;
11-MAR-1999;
                                                                                01-JUL-1999;
            23-OCT-2001
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The LS147 polypeptide is useful for preparing a composition for treating cancer, e.g. lung cancer using gene therapy. This sequence represents a universal primer used to sequence LS147 expressed sequence tag (EST)-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a purified polypeptide or its fragment derived from the LS147 gene capable of selectively hybridising to the nucleic acid of the gene and has at least 50% identity with the polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; 88; Zalphall ligand; Zalphall receptor; immune response; tumour progression; metastasis; tumour stasis; haematopoietic tumour; lymphoma; B cell tumour; systemic lupus erythematosus; rheumatoid arthritis; myaethenia gravis; diabetes; infectious disease; immunocompromised patient; HIV infection; vaccine; primer.
                                                                                                                                                                                                                                                                                                                                                         New LS147 polypeptide, useful for preparing a composition for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0
                                                                                                                                                                                                                    Klass MR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 25; DB 1; Length 26;
100.0%; Pred. No. 3.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sprecher CA, Foster DC, Holly Nelson AJ, Dillon SR, Hammond
                                                                                                                                                                                                                    Colpitts TL, Friedman PN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human cDNA encoding Zalphall sequencing primer #3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 39; 47pp; English.
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11-MAR-1999; 99US-0123904P.
01-JUL-1999; 99US-0142013P.
09-MAR-2000; 2000US-00522217.
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                                                                                                                                                                                                                    Cohen M,
Stroupe S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Presnell SR,
Johnston JV,
                                                                                                        KLASS M R.
RUSSELL J C.
STROUPE S.
                                              COLPITTS T L. FRIEDMAN P N.
BILLINGEL P.
                                                                                                                                                                                                                                                                                                  WPI; 2003-341045/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                         e.g., lung cancer
                          COHEN M.
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Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                        100.0%; Prec. ...
                                   Example 7; SEQ ID NO 39; 103pp; English.
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                                                                                                                                                                                                                                                25 AAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                exemplification of the invention.
                                                                                                                                                                                                                                                                               ADI00945 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                        LS-NOV-2002; 2002US-00295723
                                                                                                                                                                                                                                                                                                                                                                                   09-MAR-2000; 2000US-00522217
                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                          Local Similarity 100.
1es 25; Conservative
                                                                                                                                                                                                                                                                                                                                                   US2003125524-A1
                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                    22-APR-2004
                                                                                                                                                                                                                                                                                                                                                              03-JUL-2003
                                                                                                                                                                                                                                                                                         ADI00945;
                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                    RESULT 241
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The invention relates to a novel isolated zalphall ligand polypeptide. The polypeptide of the invention may be useful for boosting immunity to infectious diseases and treating immunocompromised patients, such as HIV patients, as well as in improving vaccines. The current sequence is that of the PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New zalphall ligand polynucleotide and polypeptide molecules, useful for treating cancer, e.g. melanoma, solid tumor, hematopoietic tumor, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; zalphall ligand; pharmaceutical; cancer; immune response; melanoma; tumour; solid tumour; haematopoietic tumour; lymphoma; human;
                                                                                                                                                                          New zalphall ligand polypeptides, useful for boosting immunity to infectious diseases, and treating immunocompromised patients, such as human immunodeficiency virus (HIV) patients, or in improving vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.9%; Score 25; DB 1; Length 26;
100.0%; Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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Dillon SR, Hammond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                    , Foster DC
Dillon SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human zalphall ligand PCR primer segid 39.
                                                                                                                                                                                                                                                                                  Example 7; SEQ ID NO 39; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 AAAAAAAAAAAAAAAAAAA 2733
                                                    Sprecher CA,
Nelson AJ, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sprecher CA,
Nelson AJ, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25 AAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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11-MAR-1999; 99US-0123904P.
01-JUL-1999; 99US-0142013P.
09-MAR-2000; 2000US-00522217.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25; Conservative
(ZYMO ) ZYMOGENETICS INC.
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                                                    SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SK,
                                                 Presnell (
Johnston J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Presnell (Johnston
                                                                                                                           WPI; 2003-811003/76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-440401/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR; primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-AUG-2004
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                                                 Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADP19768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                              The invention relates to stimulating an immune response in a mammal exposed to an antigen or pathogen comprises administering a composition comprising mature zalphall ligand polypeptide comprising residues 32-162 of ADH44572 in a pharmaceutical vehicle. Also included are stimulating an immune response in a mammal exposed to an antigen or pathogen comprising: (a) determining (in)directly the level of antigen or comprising zalphall ligand polypeptide in a pharmaceutical vehicle; (c) determining (in)directly the level of antigen or pathogen in the mammal; (d) comparing the antigen or pathogen level in (a) with (b), where a change in the level indicates stimulation of immune response; or pathogen comprising; (a) determining a level of antigen or pathogen. Comprising: (a) determining a level of antigen or pathogen. Comprising: (a) determining a level of antigen or pathogen. Comprising: (a) determining a level of antigen. Or pathogen. Comprising: (a) determining a level of antigen. Or pathogen. Specific antibody; (b) administering a composition comprising zalphall igand polypeptide in a pharmaceutical vehicle; (c) determining a post comparing the level of the antigen. Or pathogen-specific antibody; and (d) comparing the level of the antibody in (a) with (b), where an increase in the antibody level indicates stimulation of immune response). The method is useful for stimulating an immune response in a mammal exposed to an antigen or pathogen, and for enhancing anti-tumour activity or resulting in a reduction in tumour progression, decrease in metastasis, or tumour stasis. The tumour may be a haematopoietic tumour, a lymphoma or antigen or pathogen and is useful for treating a wide or as B cell tumour may be a haematopoietic tumour, a lymphoma or pathogen in the minimal or perhoden in the minimal or minimal 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       range of diseases arising from defects in the immune system, e.g. systemic lupus erythematosus, rheumatoid arthritis, myasthenia gravis, or diabetes, for boosting immunity to infectious diseases, treating
                                         Stimulating an immune response in a mammal exposed to an antigen or pathogen, useful for enhancing anti-tumor activity resulting in reduced tumor progression or metastasis, comprises administering zalphall ligand polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            zalphall ligand; immunity; infectious disease; immunocompromised patient; HIV; vaccine; human; ss; PCR; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunocompromised parients, such as HIV+ patients and in improving vaccines. The present sequence is a sequencing primer used in the
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100.0%; Pred. No. 3.7e+02;
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The invention describes an isolated polypeptide comprising a sequence of amino acid residues that is at least 90 or 95% identical to residues 41 (Gln) to 148 (Ile), or 32 (Gln) to 148 (Ile) of a sequence of 162 amino

Example 7; SEQ ID NO 39; 111pp; English.

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Gaps ö ngs.res

Example 7; SEQ ID NO 39; 110pp; English.

vivo, and in autologous marrow culture

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acids (SEQ ID NO:2, human zalphall ligand), fully defined in the specification. Also described are: a pharmaceutical composition composition. Also described are: a pharmaceutical composition comprising the polypeptide, and a vehicle; a method of treating cancer in a mammal; a method of stimulating an immune response in a mammal with melanoma; an stolated polymucleotide comprising a sequence of nucleotides a tumour; an isolated polymucleotide comprising a sequence of nucleotides concodes a polypeptide that binds a receptor comprising 538 amino acids, cancer a polypeptide that binds a receptor comprising 538 amino acids, fully defined in the specification; a pharmaceutical composition comprising the polymucleotide encoding, in a pharmaceutically acceptable vehicle; an expression vector comprising the following operably linked comprising and an isolated polymucleotide molecule comprising at least 10 nucleotides of the polymucleotide sequence of 642 bp, fully clean in the specification. The molecules, compositions and methods are useful for treating cancer, e.g. melanoma, solid tumour, haemacopoietic tumour, or lymphoma. This sequence represents a primer used in the composition of human cytokine zalphall ligand.
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Nelson AJ, Dillon SR, Hammond AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           stem cell; cell culture; PCR; primer; 88; zalphall ligand
                                                                                                                                                                                                                                                                                                                         Length 26;
                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human zalphall ligand-specific PCR primer - SEQ ID 39.
                                                                                                                                                                                                                                                                                             Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                  100.0%; Prec. ....
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                                                                                                                                                                                                                                                                                                                            0.9%; Score 25;
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Johnston JV,
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SPRECHER C A.
FOSTER D C.
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GROSS J A.
JOHNSTON J V.
NELSON A J.
DILLON S R.
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Matches 25; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2004260065-A1
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01-JUL-1999;
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Gross JA,
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(PRES/)
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(NELS/)
(DILL/)
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                                             The invention comprises a fusion protein that contains a zalphall ligand and a cytokine polypeptide (e.g. IL-2, IL-4, IL-15 or GM-CSF), the fusion protein of the invention binds to the human receptor protein. The protein of the invention is useful for stimulating the proliferation and/or development of hematopoietic cells. The protein of the invention is also useful in autologous marrow culture. The present DNA sequence represents a PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to novel purified polynucleotides (I; ADW14170-ADW14176) and proteins (ADW14184-ADW14187) derived from lung tissue gene LS147. The sequences are useful for detecting, diagnossing, staging, monitoring, prognosticating, in vivo imaging, preventing or treating diseases of the lung, e.g. lung cancer, pneumonia (of all origins including viral, bacterial, and fungal), asthma, black lung disease, or adult respiratory distress syndrome. Recombinant constructs comprising (I) can be produced using, e.g. plasmid pINCY, which contains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polynuclectide encoding LS147 polypeptide, useful for detecting, diagnosing, staging, preventing or treating lung diseases, e.g. lung cancer, pneumonia, asthma, or adult respiratory distress syndrome.
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0
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                                                                                                                                                                  Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                             0.9%; Score 25; DB 1; Li
100.0%; Pred. No. 3.7e+02;
ative 0; Mismatches 0;
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pe SD;
                                                                                                                                                                                                                                                                                                                                                          ADW14179 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Universal primer SEQ ID 10
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                                                                                                                                                                                                                                                                                                                                                                                                                    07-APR-2005 (first entry)
                                                                                                                                                                                                                                25; Conservative
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                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2005019820-A1
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Russell JC, S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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(COHE/)
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(KLAS/)
(RUSS/)
(STRO/)
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AAV71936 standard; DNA; 27 BP.

AAV71936/

Anchored poly T RT-PCR primer.

(first entry)

18-FEB-1999

AAV71936;

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universal priming sites adjacent to the 3' and 5' ligation junctions of the inserts. The present sequence is a universal primer used to sequence the LS147 inserts cloned in pINCY.
                                                                                                                                                                                                                                                                                                                                                                                                                            eader sequence; mRNA translation; cell-free system; protein production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New leader sequence containing a poly(A) sequence from 5A to 35A or containing an additional sequence linked to the complete or deleted construct of poly(A) sequence, useful for synthesizing polypeptides in
                                                                                                                                         Gaps
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                                                                                                    0.9%; Score 25; DB 1; Length 26; 100.0%; Pred. No. 3.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide seguence of leader seguence #1 of 5'-UTR.
                                                                   Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HOFF ) ROCHE DIAGNOSTICS GMBH.
(HOFF ) HOFFMANN LA ROCHE & CO AG F.
(PROT=) INST PROTEIN RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; SEQ ID NO 1; 26pp; English.
                                                                                                                                                                                              AEC01876 standard; DNA; 26 BP
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                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                  Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-555940/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cell-free system.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                           AEC01876;
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Matches
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The invention relates to preparation of a normalised, subdivided library of amplified cDNA from the coding regions of mRNA in a sample. The method involves reverse transcription, with at least one cDNA primer of formula convolves reverse transcription, with at least one cDNA primer of formula convolves reverse transcription, with at least one cDNA where Conl = any sequence of 1-100 nucleotides; dT = deoxythymidinyl; n2 is at least 1; n3 and n4 convolves are both 0, or n3 is 1 and n4 is at least 1; followed by second strand convolves to a similar formula, in the presence of DNA polymerase I (or its primer of a similar formula, in the presence of DNA polymerase I (or its camplification primers. Comparison of CDNA in the prepared library with a cappasse (a computer-generated list of molecular weights of restricted at abase (a computer-generated list of molecular weights of restricted convolves to known sequence) is used to detect changes in such expression (particularly for diagnosis of disease). Surfaces (chip) having amplified convolves tably immobilised on it, obtained by a similar method, are used to sorder from the family (to identify new genes) and to detect differences in the expression petterns between cells. The polypeptides expressed by the convolves in the convolves the convolves the family (to identify new genes) and to detect differences in the convolves the family (to identify new genes) and to detect differences in the convolves the con
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Preparation of normalised, subdivided cDNA libraries from mRNA - by reverse transcription and amplification, used to screen for new genes and interacting proteins, potential drugs, and for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         libraries can be used for drug development. Sequences AAV71935 to
AAV71946 represent primers used to exemplify the method of the invention
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                                                                                                                                                                                                   Normalised, cDNA library, mRNA cloning, reverse transcription, immobilise, screening, hybridisation, nucleic acid amplification, expression pattern, drug development, PCR primer, RT-PCR, 88.
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Pred. No. 3.8e+02;
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98DK-00000432.
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                                                                                                                                                                                                                                                                                                           Synthetic.
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0; Indels

0.9%; Score 25; DB 1; Length 26; 00.0%; Pred. No. 3.7e+02;

100.0%; Pred. w.

Conservative

Best Local Similarity Matches 25; Conserv

Query Match

2709 AAAAAAAAAAAAAAAAAAAAAAAA 2733

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WPI; 2002-676576/73.
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Homo sapiens.
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                                                                                                                         17-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                            Tai-Jay
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an androgen receptor complex-associated protein (ARCAP) sequence and the CDNA encoding it. The protein is useful for socreening a compound that decreases AR-mediated (androgen receptor mediated) transactivation which involves contacting the ARCAP protein with a protein complex comprising an AR in the presence of a candidate compound, measuring the extent of binding between the polypeptide, and determining if the extent of binding is less than the extent of binding between the polypeptide and the protein complex in the absence of the candidate compound. The ARCAP DNA is useful for determining if a sample contains cancerous cells which involves providing a sample from a human patient and detecting ARCAP expression in the sample. The sequences are useful for determining whether a sample contains liver tumour cells. This sequence represents a 5'RACE PCR primer used to amplify human ARCAP DNA
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                                                                                                                                                                                                                     Human androgen receptor complex-associated protein 5'RACE PCR primer #1.
                                                                                                                                                                                                                                                                             Human; androgen receptor complex-associated protein; ARCAP; primer; ss; androgen receptor; AR; cancer; liver tumour; cytostatic; PCR; 5'RACE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel substantially pure androgen receptor (AR) complex-associated protein which binds to AR and increases ability of AR to transactivate androgen-responsive gene, useful as drug target for treating liver
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                                BP.
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12-FEB-2001; 2001US-00781693.
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                                ABS53863 standard; DNA; 27
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Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1227150-A2.
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      ABS53863/c
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The present invention relates to the isolation of human androgen receptor complex-coupled protein (ARCAP), and the polynuclectide sequence encoding it. The ARCAP polypeptide complexes with an androgen receptor to increase the activity of the androgen receptor, transactivating the androgen responsing gene. The invention also describes a a vector containing the ARCAP polynuclectide sequence, and a host call containing the ARCAP polynuclectide sequence. The ARCAP polypeptide can be used as a treating agent. The present sequence represents a PCR primer used in the example of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                 Novel substantially pure androgen receptor (AR) complex-associated protein which binds to AR and increases ability of AR to transactivate androgen-responsive gene, useful as drug target for treating liver
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100.0%; Pred. No. 3.8e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                              (VETE-) VETERANS GEN HOSPITAL
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28-FEB-2001; 2001JP-00055192.
                                                                        12-FEB-2001; 2001US-00781693
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nes 25; Conservative
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WPI; 2002-437469/47
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modified_base
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                                                                                           Multivalent
specificity
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                                               Walker SM;
                                                                                                                              disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 251
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                                                                                                                                                                                           The invention relates to a novel multivalent compound comprising two binding groups specific for different binding sites on the same target. The compound of the invention demonstrates cytostatic, antitheumatic, antiatrchritic, antiporatic, antidiabetic, ophthalmological, antiatrchritic, antiulcer and vasotropic activities and may act as an inhibitor of receptor tyrosine kinase activity. The compound may be used to prepare diagnostic imaging agents and pharmaceutical compositions for treating diseases associated with angiogenesis or hyperproliferation, particularly tunmours, but also theumatoid arthritis, psoriasis, diabetic retinopathy, atherosclerosis, ulcers and restenosis. Furthermore, the compound may be utilised as a contraceptive via inhibition of uterine neovascullarisation. The current sequence is that of the RT-PCR primer region)-related RNA.
                                                                                                                               as
                                                         Ramalingam K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; primer; antiarthritic; cytostatic; ophthalmological; angiogenesis inhibitor; Kdr tyrosine kinase inhibitor; VEGF antagonist; hepatcocyte growth factor antagonist; multivalent compound; binding moiety; euplastic tumour growth; angiogenesis; hyperproliferation; arthritis; atherosclerotic plaque; corneal graft neovascularization; ocular disease.
                                                                                                                          New compound with two different binding groups for same target, useful diagnostic and therapeutic agent, e.g. for tumors and other angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                           Arbogast C, Bussat P, Dransfield DT, Fan H, Linder KE;
Marinelli ER, Nanjappan P, Nunn A, Pillai R, Pochon S, Ra
Sato A, Shrivastava A, Song B, Swenson RE, Von Wronski MA;
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                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 25; DB 1; Length 27; 100.0%; Pred. No. 3.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                            Sequence 27 BP; 0 A; 1 C; 1 G; 25 T; 0 U; 0 Other;
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                                                                                                                                                                       Example 6; Page 119; 278pp; English.
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            В.
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                                                                                                   WPI; 2004-053022/05.
            BRACCO INT
                        DYAX CORP
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                                                                            Walker SM;
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                                                                                                                                                   diseases.
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          (BRAC)
                        (DYAX-)
                                                                  Sato A,
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                                     Ramalingam K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a multivalent compound (C) comprising at least two binding moieties having specificity for different binding sites on the same target. (C) is useful for treating euplastic tumour growth and disease associated with angiogenesis or hyperproliferation (claimed). (C) is useful for treating diseases such as arthritis, atherosclerotic is useful for treating diseases such as arthritis, atherosclerotic bladues, corneal graft neovascularization or ocular diseases. (C) is small and can more easily reach a target. (C) localizes more effectively to the target site than other targeting compounds due to its binding to more than one site on the same target. This sequence represents a DNA more than one site on the same target.
                                                                                                                                                                                                                                Multivalent compounds with at least two binding moities having specificity for different binding sites on the same target, useful for treating and diagnosing, e.g. angiogenic and hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Arbogast C, Bussat P, Dransfield DT, Fan H, Linder K;
Marinelli ER, Nanjappan P, Nunn A, Pillai R, Pochon S, Ra
Sato A, Shrivastava A, Song B, Swenson RE, Von Wronski MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 25; DB 1; Length 27; 100.0%; Pred. No. 3.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                              Example 6; SEQ ID NO 72; 320pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide used in the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100.
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                                                                                                                                                                  WPI; 2004-593275/57
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The present invention relates to a method for analysing a target nucleic acid consisting of repetitive and non-repetitive sequences. The method comprises hybridising a polynucleotide probe comprising a segment complementary to a non-repetitive region and a segment complementary to an on-repetitive region and a segment complementary of an adjacent repetitive region, where the second segment consists of a defined number of repeats, and determining the melting point temperature of the hybrid. The method is used to analyse microsatellites, especially microsatellite instability, particularly as a means for detecting meredicaty tumours. Alternatively, the method is used to identify an individual in a population. The present sequence is a probe for Mononucleotide repeat locus BAT25, and was used to illustrate the
Analyzing repeat sequences in DNA using a probe which hybridizes to adjacent repetitive and non-repetitive regions and determining hybrid melting point is useful to detect microsatellite instability such as in hereditary cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29 BP; 26 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Prec.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.9%; Score 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 AAAAAAAAAAAAAAAAAAAAA 26
                                                                                                                          English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match
Best Local Similarity 100.
Matches 25; Conservative
                                                                                                                             Claim 16; Page 7; 19pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                        invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADO81068;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 252
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Gaps

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DB 1; Length 29; 4e+02; hes 0; Indels

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gene typing; polymorphic microsatellite loci; PML; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pTT-blue-vector; cow; microsatellite; PCR; primer; ss.
                                                  Cow prion protein microsatellite locus primer #80.
 ВР
                                                                                                                                                                         09-AUG-2002; 2002DE-01036711.
ADO81068 standard; DNA; 28
                                  (first entry)
                                                                                                                                       DE10236711-A1
                                                                                                                                                        26-FEB-2004.
                                                                                                                      Bos taurus
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The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons. Example 3; Page 28; 64pp; German. 09-AUG-2002; 2002DE-01036711. ģ (UYHO-) UNIV HOHENHEIM. Geldermann H, Preuss WPI; 2004-215730/21.

Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; adrenal gland disorder;

Transgenic mouse; neurological disorder; ardiovascular disorder;

Transgenic metabolic disorder; immune disorder; bone disorder;

Transgenic metabolic disorder; nutritive disorder; cancer;

Transgenic metabolic disorder; nutritive disorder; cancer;

Transgenic metabolic disorder; prostate disorder; cancer;

Transgenic metabolic disorder; prostate disorder; pleen disorder;

Transgenic metabolic disorder; pancreas disorder; spleen disorder;

Transgenic metabolic disorder; pancreas disorder; spleen disorder;

Transgenic metabolic disorder; pancreas disorder; spleen disorder;

Transgenic metabolic metabolic; antiparkinsonian; antimanic;

Transgenic metabolic metabolic metabolic; antidiabetic;

Transgenic metabolic metabolic metabolic; antidiabetic;

Transgenic metabolic metabolic metabolic metabolic;

Transgenic metabolic metabo determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and predisposition (M3) of diseases associated with gene that include bML. The method is used to identify microsatellite markers, in a disease-related gene, that are associated with a predisposition to diseases and for prediagnosis of such diseases, aspecially prion to but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is sinpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus. Gaps Novel mammalian G protein coupled receptors, useful for identifying compounds that modulates diagnosing and treating disease condition associated with GPCR dysfunction e.g. autoimmune diseases, angina pectoris, Parkinson's disease. ö 0.9%; Score 24.8; DB 1; Length 28; 92.9%; Pred. No. 4e+02; Indels Hohmann J, JE, Gragerov A, Hohmann J, Pavlova MN, Vassilatis D, Sequence 28 BP; 0 A; 2 C; 0 G; 26 T; 0 U; 0 Other; ; 0; Mismatches 2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2736 primer, SEQ ID NO:1598. 09-SEP-2002; 2002US-0409303P. 09-SEP-2003; 2003WO-US028226 Gaitanaris GA, Bergmann JE, Madisen L, Mcilwain KL, Pa ADO30495 standard; DNA; 26 29-JUL-2004 (first entry) 26; Conservative WPI; 2004-390329/36. (PRIM-) PRIMAL INC. Best Local Similarity WO20040400000-A2 13-MAY-2004. 5' RACE PCR Madisen L, Synthetic. ADO30495; Query Match Matches RESULT 253 AD030495/c 888888888888888888 THE STANKE STREET STREE g ઠે

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The invention relates to human and mouse G protein-coupled receptors

(GPCRS) and nucleic acids encoding them. The invention also relates to

sequences at least 90% identical to the GPCR proteins and nucleic acids

of the invention; methods of treating, preventing or diagnosing diseases

consounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR-related diseases; a transgenic

compounds useful in the treatment of GPCR per invention; and wite coupling decreases

compounds and different GPCR polynucleotides of the invention. The

compounds useful in the discloses variants of the GPCR polypeptides and vectors

comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may

be used in the diagnosis, treatment or prevention of a wide variety of

diseases including neurological disorders (e.g., Alzheimer's disease,

depression, diabetic neuropathy, Parkinson's disease or schizophrenia);

compounds and disorders of the colon or intestine

compounds and disorders of the colon or intestine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Syndrome); cardiovascular disorders (e.g., angina, cardiac arrhythmia or myocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or myocardial infarction); muscular disorders (e.g., autoimmune disorders (e.g., antoimmune disorders (e.g., antoimmune disorders (e.g., antoimmune disorders or AIDS); bone and joint disorders (e.g., osteoarthritis, rheumatoid arrhitis, gout or osteoporosis); metabolic or nutritive disorders (e.g., obesity, enzyme deficiency-related diseases or vitamin deficiency-related diseases); and disorders of the kidney, liver, lung, breast, ovary, uterus, prostate, testis, skin, stomach, pancreas, spleen, thymus and thyroid (e.g., cancers). The present sequence represents a RACE (rapid mylification of cDNA ends) PCR primer used in the isolation of cDNA encoding human GPCRs. Note: The full sequence data for this patent did not form part of the printed specification; those sequences not shown the contained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inhibitor; hyperproliferative disease; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POLYA, a competitor oligonucleotide for binding human PUR-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 24.2; DB 1; Length 26; 96.0%; Pred. No. 4.2e+02; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 2 Other;
                        Disclosure; SEQ ID NO 1598; 542pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ftp.wipo.int/pub/published_pct_sequences.
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93US-00014943.
95US-00470911.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT99286 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PUR element; human; c-myc; in
cancer; probe; hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00486421
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Best Local Similarity 96.vv
Best Local 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-FEB-1993;
06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5672479-A
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Gaps

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                                                                                                                                                                The oligonucleotides AAT99279-T99286 were used as competitor digonucleotides for the binding of PUR pritein to DNA. The PUR sequence can be used to identify chemical or biological compounds that bind to PUR or binding fragments of PUR. Inhibitors of PUR activity may be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This is the nucleotide sequence of an oligonucleotide used as a competitor with the PUR element in the method of the invention, involving the use of the PUR protein and its fragments, which inhibit PUR protein binding to PUR element or other proteins. Inhibitors of PUR activity may be useful for treating viral infections and hyperproliferative diseases
                                                                                         Assays for PUR protein ligands or modulators - using immobilised PUR protein or fragments, to treat hyper-proliferative diseases, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein and its fragments - that inhibit PUR protein binding to PUR
                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibition; viral infection; cancer; PUR element; ive disease; ss.
                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                         0.9%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 4.2e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 24;
                                                                                                                                                                                                                                                            Seguence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of the oligonucleotide POLYA.
                                                                                                                                                                                                                                treat hyperproliferative diseases such as cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 24;
                                                                                                                                                                                                                                                                                                                                                      2709 AAAAAAAAAAAAAAAAAAA 2732
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                                                                                                                                                                                                                                                                                                                                                                                  1 AAAAAAAAAAAAAAAAAAAAA 24
(MOUN ) MOUNT SINAI SCHOOL MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MOUN ) MOUNT SINAI SCHOOL MEDICINE.
                                                                                                                                    Example; Col 33; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
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93US-00014943.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PUR-alpha gene; inhibition;
hyperproliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV31743 standard; DNA; 24
                              Bergemann AD, Johnson EM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              element or other proteins
                                                                                                                                                                                                                                                                                                                       24; Conservative
                                                          WPI; 1997-488859/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-321632/28
                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   such as cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 255
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pBluescriptSK+ phagemid primer SEQ ID NO:
                                                                                                                                                   Romantchikov Y;
                                                       WO200036088-A1
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                                                                                                              17-DEC-1998;
                                                                                           17-DEC-1999;
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                                                                         22-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                      Synthetic
                                                                                                                                                                                                                    libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA40353;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 258
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                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a monoclonal antibody that specifically binds to an epitope of the PUR protein. Antibodies that bind to the PUR protein and neutralise PUR activity may be used to treat hyperproliferative diseases such as cancer. PUR antibodies may be used diagnostically to detect aberrant expression of the PUR protein and/or matations in the PUR gene. The present sequence represents an oligonuclectide used in the cloning and sequencing of the PUR protein and its sequence element PUR repeat, in an example from the present invention
          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                 Monoclonal antibody specific for PUR protein - useful for treating
                                                                                                                                                                      PUR element, PUR-alpha, hyperproliferative disease, cancer, human, monoclonal antibody, identification, characterisation, 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.9%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 4.2e+02;
ative 0; Mismatches 0; Indels
                                                                                                                                                   Oligonucleotide POLYA used in PUR cloning and sequencing.
          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
 Pred. No. 4.2e+02; Mismatches 0;
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                             2709 AAAAAAAAAAAAAAAAAAA 2732
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                                               1 AAAAAAAAAAAAAAAAAAAAAAAAAA 24
                                                                                                                                                                                                                                                                                                                          SINAI SCHOOL MEDICINE.
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                                                                                                                                                                                                                                                                                                                                                                                                              Example; Col 33; 64pp; English.
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93US-00014943.
95US-00470911.
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                                                                                              AAX04086 standard; DNA; 24 BP
 100.08;
                                                                                                                                                                                                                                                                   95US-00486809
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                                                                                                                                   12-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                              Johnson EM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24; Conservative
            24; Conservative
                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-152881/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
  Best Local Similarity
                                                                                                                                                                                                                                                                                                                           TNUOM ( NUOM)
                                                                                                                                                                                                                                                                                                                                               Bergemann AD,
                                                                                                                                                                                                                                                                                     28-AUG-1992;
02-FEB-1993;
06-JUN-1995;
                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                    07-JUN-1995;
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                                                                                                                                                                                                                                                 09-FEB-1999
                                                                                                                                                                                                     Synthetic
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                                                                                                                 AAX04086;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                               cancer.
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                                                                             RESULT 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
             Matches
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This invention describes a novel method (M1) for inserting a nucleic acid (N1) into a circular vector (V1) comprising joining ends of N1 and V1 under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reannealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex populations of DNA or DNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger nucleic acid fragments are just as easily cloned, allowing highly representative libraries to be made. Vector to vector ligation is avoided using the methods. AAA4031-A40366 represents primers used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inserting a nucleic acid into a circular vector comprising joining their ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
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Primer; cloning; ligation; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ROMA/) ROMANTCHIKOV Y
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                                                                                                   Inserting a nucleic acid into a circular vector comprising joining their
                                                                                                                 ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaccinating against tumors, infectious diseases, allergies and asthmausing immunostimulatory Py-rich and TG nucleic acids.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #872.
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                                                                                                                                                                                   Example 1; Page 66; 71pp; English
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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                                                                                                                                                                                                                                                                                                                                                                                                                the method of the invention
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(ROMA/) ROMANTCHIKOV Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Krieg AM, Schetter C,
                                                                 WPI; 2000-442381/38
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                                   Romantchikov Y;
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                                                                                                                                                       libraries
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                                                              response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immuno response. The present sequence is one such immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects adainst tumour antigens, viral antigens (e.g. herpesviridae, retroviridae, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or
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                                                                                                                                                                                                                                                    immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
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                                                    The present invention relates to a method for stimulating an immune
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                                                                                                                                                                                                                                                                                            sequence may have a phosphorothioate backbone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory nucleic acid #420.
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                  Claim 101; Page 57; 338pp; English.
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(COLE-) COLEY PHARM GMBH.
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                 against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Bacherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or fimmune deficiency. The present sequence an also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
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py-rich) or thymidine (T) rich. The method is used to vaccinate subjects
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using immunostimulatory Py-rich and TG nucleic acids.
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4.2e+02;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; prostate cancer; cytostatic; carcinogen; pharmacodyanamic marker; pharmacogenomic marker; gene; ss.
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                                                                                                            Length 24;
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                                                                                                                                                                   0; Indels
present sequence may have a phosphorothioate backbone
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                                                      Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                          0.9%; Score 24; DB 1; Le
100.0%; Pred. No. 4.2e+02;
rative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human prostate expression marker cDNA 14833.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
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                                                                                                                                                                                                                                                         Monahan JE;
                                                                                                                                                                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-FEB-2000; 2000US-0183319P.
16-WAR-2000; 2000US-0189862P.
25-MAY-2000; 2000US-0207454P.
09-JUN-2000; 2000US-021314P.
18-JUL-2000; 2000US-0219007P.
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                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                         24;
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                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                      RESULT 262
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(first entry)

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Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                       Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection, neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                Angiogenesis inhibitory oligonucleotide #433.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; Page 27; 276pp; English
                                                                                                                                                                                                                                                                                                                 scleroderma, hypertrophic scar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (COLE-) COLEY PHARM GROUP INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-DEC-2001; 2001WO-US048458.
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                                                                       13-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUL-2002,
                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                             ABS77949;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumacoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                        Angiogenesis inhibitor; se; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; heemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to inhibiting angiogenesis in a subject,
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Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                              Angiogenesis inhibitory oligonucleotide #961.
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100.0%; Pred. No. ...
                         AAAAAAAAAAAAAAAAAAAAAA
Claim 2; Page 36; 276pp; English
                                                                                                                                     ВР
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                                                                                                                              ABS78477 standard; DNA; 24
                                                                                                                                                                                                                       (first entry)
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Best Local Similarity 100.
Matches 24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                            ABS78477;
                         24
                                                                                                              ABS78477
                                                                                         RESULT
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The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, obler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, intestinal adhesions, atherosclerosis, scleroderma and wound granulation, intestinal adhesions, atherosclerosis, scleroderma and
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                                                                                                                                                                                                                                                                                                                            hypertrophic scars. The present sequence is an antiangiogenic nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indela
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2709 AAAAAAAAAAAAAAAAAAAA 2732
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS78478 standard; DNA;
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Gaps .; 0

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ABS77949 standard; DNA; 24 BP.

RESULT 264 ABS77949/c ID ABS77

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia;
                                                                      rubeosis; Osler Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
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Pred. No. 4.2e+02;
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100.0%; Pred. No...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 36; 276pp; English.
                                                                                                                             scleroderma; hypertrophic scar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABL39405 standard; DNA; 24 BP
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                                                                                                                                                                                                                                                                      14-DEC-2001; 2001WO-US048458.
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les 24; Conserv
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                Bratzler RL;
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                                                                                                                                                                 Synthetic.
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The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigons and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, concer, leukaemia, liver cancer, elegencer, Hodgkin's lymphoma, non-CC cancer, pancreatic cancer, myeloma, oral cavity cancer, ovarian CC cancer, pancreatic cancer, prostate cancer, indexoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The CC cancer, stomach cancer testicular cancer, and uterine cancer. The concernation of the invention
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                                                                                                                                                                                                                                                                                       Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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/*tag= a
/mod_base= OTHER
/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.9%; Score 24; DB 1; L
100.0%; Pred. No. 4.2e+02;
ive 0; Mismatches 0;
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                                                                                                                                                               22-JUN-2000; 2000US-0213346P.
                                                                                                                              22-JUN-2001; 2001WO-US020154
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Matches 24; Conservative
                                                                                                                                                                                                                             Hartmann G;
                                                                                                                                                                                                                                                             WPI; 2002-154611/20.
                                                                                                                                                                                                                                                                                                                                             developing cancer
                                                                  WO200197843-A2
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modified_base
                                                                                                  27-DEC-2001.
                                                                                                                                                                                                                             Weiner G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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Polymorphism detection; sequence detection; mutation detection; A24; probe; non-dissociative termolecular complex; dopTAR sensitiser particle; single nucleotide polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A24 oligonucleotide used to create dopTAR chemiluminescer particles.
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/note= "A is covalently linked to a
PO2OCH2CH2CH2CH2CH2OH moiety"
                                                                                                                                                                                                                                                                                                                                   100.0%; Pred. No. 4.2e+02;
tive 0; Mismatches 0;
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                                                                                                                                            Example; Page 58; 87pp; English
                                                                          Patel R,
                                                                                                                                                                                                                                                                                                                                                                                                                      AAS17869 standard; DNA; 24 BP.
                        03-MAY-2001; 2001WO-US014528.
                                        04-MAY-2000; 2000US-00564230
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                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Marches 24; Conservative
                                                         (DADE-) DADE BEHRING INC.
                                                                         Pease JS, Cromer R,
                                                                                          WPI; 2002-164078/21.
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       08-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS17869;
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 268
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This invention represents a method for detecting the presence of a polymucleotide sequence, differences in polymucleotide sequences or mutations in genomic DNA. The method involves contacting 3 coligonucleotide probes with a sample contacting a polymucleotide. The clirat probe hybridises to a region of the polymucleotide. The second and third probes can bind a second region of the polymucleotide contacting a second and third probes are identical except for the sequence. The second and third probes are identical except for the sequence or difference of one or more nucleotides. The reaction medium is presence or difference of one or more nucleotides. The reaction medium is contacted to conditions for forming substantially non-dissociative termolecular complexes, which can be at least one of, the polymucleotide sequence with the first and third probes. The oligonucleotide probes have labels on covalently bound to allow for their detection upon binding. The method of the invention is useful for detecting the presence of a single concovalent properties of a single concovalent properties (SNP) in a fragment of genomic DNA. The method can be used for the direct detection of nucleic acid in very small contacting amplification. In addition, the method may be carried contacting amplification of the target and reference sequences.
                                                                                                                                                                                                                                                                                                                             Detecting presence of polynucleotide, differences between polynucleotide sequences, useful for detecting single nucleotide polymorphism and alleles of polynucleotide sequence involves use of three competitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence represents an oligonuclectide probe A24 used to create dopTAR chimiluminescer sensitiser particles in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Binding the nucleic acid to a suspendable particle acts as a support provides a means of segregating the bound polynucleotide target from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 47; 75pp; English.
   17-MAY-2001; 2001WO-US016089.
                                                                                                                                   (DADE-) DADE BEHRING INC
                                                                                                                                                                                                                                                                  WPI; 2002-097664/13.
                                                              19-MAY-2000;
                                                                                                                                                                                                 Patel RD;
                                                                                                                                                                                                                                                                                                                                                                                                                                 probes.
The invention relates to the detection of multiple components in a medium, comprising combining the medium with at least two sensitiser reagents, and at least one reactive reagent activated by a product can entire the sensitiser reagents when activated; and differentially activating the sensitiser reagents. The combination of sensitiser reagents and reactive reagent (s) allows differential detection of the components. Methods of the invention may be used for the detection of ligands, receptors and polynucleotides, and also for the detection of components, Mosphate esters, thiophosphates, carbamates and polyhalogenated sulfenamides) and pollutants. Methods of the invention allow the detection of multiple analytes in a single test medium. An application of the methods of the present invention would be in the field of clinical diagnostics. The current sequence represents A24 oligonucleotide for the creation of oligonucleotide coated phthalocyanine sensitiser particles (PC-A24)
                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of multiple analytes, e.g. ligands, receptors, polynucleotides and pollutants, involves adding a combination of sensitizer reagents and reactive reagent Actuatable by a product of the sensitizer reagents.
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0.9%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 4.2e+02; 0; Indels Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 1 Similarity 100.0%; Pred. No. 4.2 24; Conservative 0; Mismatches 2709 AAAAAAAAAAAAAAAAAA 2732 1 AAAAAAAAAAAAAAAAAAAAAA 24 Local Similarity Query Match Matches ABK15639, 8 g

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Gaps

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0; Indels

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Gaps ö

> ABK15639 standard; DNA; 24 BP. (first entry) 08-MAY-2002 ABK15639;

RNA-PCR; primer; ss; poly(dT)24; cytostatic; antibacterial; gene therapy; manka-cDNA hybrid; gene function inhibition; cancer; PTGS; antiBense; high throughput screening; D-NNA; DNA-RNA interference; RGRp; RNA dependent RNA polymerase; posttranscriptional gene silencing. RNA-PCR procedure primer poly(dT)24.

WO200210374-A2 Synthetic 

07-FEB-2002

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The invention relates to generating mRNA-cDNA hybrids, comprising (a) providing a solution containing a nucleic acid template, one or more primars complementary to the sense conformation of the nucleic acid template, and with an RNA cemplate, and one or more promoter-linked primars complementary to the antiesnse conformation of the nucleic acid template, and with an RNA prime to synthesise a first cDNA strand, (c) treating the first cDNA certain with one or more promoter-linked primers to synthesise a promoter-linked double-stranded nucleic acid, (d) treating the promoter-linked double-stranded nucleic acid, (d) treating the promoter-linked couble-stranded nucleic acid, (d) treating the promoter-linked couble-stranded nucleic acid, (d) treating the promoter-linked couble-stranded nucleic acid, (e) treating the mRNA fragments with one or more primers to synthesise and (e) treating the mRNA fragments with one or more primers of pure and consult of the amplified mRNA.

CC double-stranded nucleic acid to synthesise amplified mRNA fragments and secretific mRNA-cDNA hybrids for transducing biological effects of interest in vitro as well as in vivo, for inhibiting gene function in prokaryotes and eukaryotes in vivo and in vitro, for suppressing cancer-related condidate molecular pathways with systematic knock out of involved molecules, in high throughput screening of gene functions in studying candidate molecular pathways with systematic knock out of involved molecules, in high throughput screening of gene function in the physiological conditions. The mRNA-cDNA hybrids may be used to screen for special gene functions, for manipulating gene expression in vitro, and comporation to increase the stability and effectiveness of transfected provide higher affinity of the mRNA hybrids may burdleotide analogue conditions. The mRNA comparase) enzyme may due to lower hinding interaction between NNA-NNA winderses harm RNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA due to lower binding interaction between DNA-RNA duplexes than RNA-RNA duplexes. The cDNA part of a D-RNAi provides further antisense gene knockout activity in addition to the posttranscriptional gene silencing (PTGS) mechanisms of the sense-RNA template, resulting in multiple specific gene interference effects with one probe. The present sequence is a poly(dT) PCR primer used in conjunction with oligo(dC)loN primers to reverse transcribe mRNA into first strand cDNA in the method of the
                                                                                                                                                                                                                                            Generating mRNA-cDNA hybrids for suppressing cancer-related genes, or treating or preventing microbe related genes, comprises thermocycling steps of promoter-linked double-stranded cDNA or RNA synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                   Example 5; Page 26; 53pp; English
                                                                                                    (UYSC-) UNIV SOUTHERN CALIFORNIA.
                                                                                                                                                Chuong C, Widelitz RB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ80181 standard; DNA; 24 BP
02-AUG-2001; 2001WO-US024412.
                                                02-AUG-2000; 2000US-0222479P.
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Immunostimulatory oligonucleotide SEQ ID NO:53

(first entry)

23-MAY-2003 ABZ80181;

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The present invention describes immunostimulatory nucleic acids of 14-100 nucleotides in length comprising the formula 5' XIDCGHX2 3' (I), where XI or X2 = independently any sequence 0-10 nucleotides, D = nucleotide other CC than C; C = cytosine; G = guanine; H = nucleotide other than G: The cimmunostimulatory nucleic acid further comprises a sequence consisting of immunostimulatory nucleic acid further comprises a sequence consisting of P and N positioned immediately 5' to X1 or 3' to X2 and N is a B cell contralising sequence, where N begins with a CGG trinucleotide and is at contralising sequence, where N begins with a CGG trinucleotide and is at cleast 10 nucleotides long. Also described: (1) a pharmaceutical composition comprising the immunostimulatory nucleic acid and a carrier; cand (2) treating an allergic condition. (I) has antiallergic activity and can be used in gene therapy. (I) can be used for preparing a composition of for treating a variety of immune related disorders such as cancer. CC infectious diseases and allergic disorders. (I) also stimulates the cativation of natural killer cells and the production of type 1 cc interferon (IFN). The present sequence represents an immunostimulatory composition composition and the present invention composition composition and carrier colligonucleotide, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New immunostimulatory nucleic acid, useful for preparing a composition for treating an allergic condition.
Immunostimulation; immune response; natural killer cell; interferon; type 1 interferon; IFN; cancer; infectious disease; allergic disorder; immune related disorder; ss.
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                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                   (COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH. (IOWA ) UNIV IOWA RES FOUND.
                                                                                                                                                                                                                                                                                                                         19-AUG-2002; 2002WO-US026468.
                                                                                                                                                                                                                                                                                                                                                              2001US-0313273P.
2002US-0393952P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vollmer J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-268241/26.
                                                                                                                                                                                                                                             WO2003015711-A2.
                                                                                                                          Key
modified_base
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03-JUL-2002;
                                                                                                                                                                                                                                                                                27-FEB-2003
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                                                                                   Synthetic
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      2X2X5X8X
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Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;

88; PCR; RT-PCR; primer; reverse transcriptase PCR; antisense therapy; mRNA expression profile; promoter containing primer.

Synthetic.

US2003022318-A1.

30-JAN-2003.

07-SEP-2001; 2001US-00949305.

25-JAN-2000; 2000US-00494212

(EPIC-) EPICLONE INC.

Lin S, Ying S;

WPI; 2003-479488/45.

Improved polymerase thermocycling reaction for nucleic acid amplification, by thermal cycling of promoter-linked nucleic acid replate synthesis and in vitro transcriptional amplification of nucleic

Example 7; Page 14; 28pp; English.

The invention relates to an improved polymerase thermocycling reaction (MI) for linear amplification of nucleic acid templates (I), combining the denaturing a number of nucleic acid templates (I), combining the denaturing a number of parimer (P2), a number of deoxynucleotide triphosphates and ribonucleotide triphosphates and muber of deoxynucleotide triphosphates and ribonucleotide triphosphates and muber of deoxynucleotide triphosphates and ribonucleotide triphosphates.

CC ontacting templates, denaturing the promoter-containing templates to contacting the denatured promoter-containing templates of promoter-containing templates, denatured promoter-containing templates, denatured promoter-containing double-stranded mucleic acid templates are flamated by Pi in one and and P2 in the other and of the other orientation, transcribing the promoter-containing double-stranded bub templates. Conteating the primer region of the promoter-containing double-stranded bub templates, conteating the primer region of the promoter-containing double-stranded bub templates, conteating the primer region of the primer region of the primer amplified RNA sequences with P2 to form a number of cDNA and a number of promoter-containing double-stranded by templates, conteating double-stranded by templates, conteating double-stranded by templates, conteating double-stranded by templates, conteating double-stranded bolymerase thermocycling reaction for linear amplification of mucleic acid sequences, and thus for producing mRNA expression profile of a cell by M1 to generate multiple copies of the calls in both normal and diseased state, amplifying the diseased state, by generating aberrant protein production of the protein coded for by the isolating aberrant protein production of the protein coded for by the isolating aberrant protein production of the protein coded for by the isolated mRNA. M1 is also useful for treating a percent expression of a cell in a diseased state caused by behavior deservation production of cell in a diseased state ca

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The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                      Gaps
                                                                                                                                                                                                                                                                                                                                   Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine; non-allergic inflammatory disease, psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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   Length 24;
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100.0%; Pred. No. 4.2e+02;
                                  0; Indels
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 Score 24; DB 1; Lo
Pred. No. 4.2e+02;
Query Match 0.9%; Score 24; DB Best Local Similarity 100.0%; Pred. No. 4.2 Matches 24; Conservative 0; Mismatches
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Best Local Similarity 100.0%; Pred. No. 4.4
Conservative 0; Mismatches
                                                                     2709 AAAAAAAAAAAAAAAAAAA 2732
                                                                                                                                                                                                                                                                                                  Immunostimulatory nucleic acid #415.
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                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           JS2003050268-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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/*tag= a
/note= "Binds to nucleotides 42-19 of the 1st strand cDNA
                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                  Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; nucleic acid amplífication; multiple step elimination; varying reaction condition elimination; poly A tract.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 24;
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Local Similarity 100.0%; Score 24; DB 1; Length 24;
Local Similarity 100.0%; Pred. No. 4.2e+02;
hes 24; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 34; 229pp; English.
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                                     29-MAR-2002; 2002US-00112653.
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                                                                                                                                                                                                                               WPI; 2003-521815/49
                                                                                                                                                                                       Krieg AM, Berg DJ;
                                                                                                                         KRIE/) KRIEG A M.
                                                                                                                                             (BERG/) BERG D J.
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13-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatisis, latex dermatitis or inflammatory bowel disease e.g., ulcerative collitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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                                                                                                                       Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine; non-allergic inflammatory disease; psoriasis; eczema, allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colltis; Crohn's disease; ss.
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                                                                                 Immunostimulatory nucleic acid #920.
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Matches 24; Conservative
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Gaps

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Comprises primer extension by reverse transcriptase and hybridising an oligonucleotide to the single stranded DNA, where the oligonucleotide is blocked from extension at the 3' end. The method is useful for amplification of nucleic acids. In the new method, a promoter is protected from degradation throughout the method. The promoter is constructed so that it does not serve as a primer for extension of a sequence that is complementary to the target sequence, i.e. it is blocked. The method can be combined with other processes to eliminate the need for multiple steps and varying reaction conditions and their associated problems. At least three otherwise separate enzymatic reactions can occur consecutively in one phase (i.e., without organic extraction and precipitation), more preferably in the same reaction vessel. Preferably, cDNA synthesis according to the new method may occur in a modifical low salt buffer. The present sequence represents the poly A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                          invention relates to a method of amplification of nucleic acid which
                 Amplification of nucleic acids, where the promoter is blocked from extension at the 3' end, useful for eliminating multiple step reactions
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                                                                                                                                                                                                                                                                                                                                                                                     tract of a mRNA used to illustrate the method of the invention.
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Pred. No. 4.2e+02;
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                                                                        Disclosure, Fig 2, 9pp, English
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Best Local Similarity 100.(
Matches 24; Conservative
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PETERSEN D
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invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
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                                                                                     0.9%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 4.2e+02;
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(PETE/) PETERSEN D
                                                                                                    Similarity
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The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-respondive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
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                                                                             ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
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Best Local Similarity 100.0%; Fred. No. 4.2
Marches 24; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                Fouron Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory nucleic acid #873.
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(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
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                             04-DEC-2003
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                                                                                                                          Synthetic.
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ADB37259;
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The present invention describes a merical to the combination products between two nucleotide sequences. The method comprises combining an initial set of oligonucleotides corresponding to a distinct nucleotide sequence and the authorities and further combining the corresponding to a distinct nucleotide sequence and further combining the initial and subsequent sets of combination oligonucleotides having a sequence region corresponding to the initial nucleotide sequence and a sequence region corresponding to the second oligonucleotide sequence. Also described is a method of creating a collection of recombination products between two genes. The methods and compositions of the present invention are useful in the field of synthetic gene technology, and more specifically, to generating a collection of recombination products between distinct nucleotide sequences. They can also be used in cassembling a library, or a population or a collection of polypeptide variants that correspond to single or multiple polynucleotide recombination products. The present sequence is used in the cerombination of the present invention.
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                                                                                                                                                                                                                           The present invention describes a method for creating a collection of recombination products between two nucleotide sequences. The method
                                                                                                Creating recombination products between two distinct nucleotide sequences, useful in the field of synthetic gene technology, and in assembling a library, or a population or a collection of polypeptide
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.00.0%; Pred. No. 4.2e+02;
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/note== "optional biotin label"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Prec. ...
                                                                                                                                                                                          Example 3; SEQ ID NO 106; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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/mod_base= OTHER
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(EGEA-) EGEA BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE25524 standard; DNA; 24
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                                                                 WPI; 2003-663477/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
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modified_base
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                                    Evans GA;
                                                                                                                                                           variants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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WPI; 2003-697450/66
Xia J;
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Detecting nucleic acid targets, useful e.g. for diagnosing single nucleotide polymorphisms, by extension of capture probe complementary to open circle probe.

Example 1; Fig 5; 66pp; English.

The invention is directed to novel methods of amplifying and detecting CC DNA using rolling circle amplification (RCA). The invention relates to detecting a target sequence (I), which involves using a capture probe (CC DNA that is complementary to an open circle probe and includes a capture probe (CP) that is complementary to an open circle probe and includes one domain complementary to an OCP (Open circle probe) and a second domain that contains a CC cleavage site (CS), to form a device; treating CP with (I) and OCP for form a hybridisation complex (HC); treating CP with a ligase so that OCP is circularised, forming a second complex (HC); treating CP with a C cleavage agent, to cut at CS, and adding an extended CP, which is cleavage agent, to cut at CS, and adding an extension enzyme (ES) and cleavage agent, to cut at CS, and adding an extension enzyme (ES) and cleavage agent, to cut at CS, and adding an extension enzyme (ES) and cloading is used for detection, genotyping and/or quantification of the method is used for detection, genotyping and/or quantification of the esting, particularly detection of single-nucleotide polymorphisms. The method permits a high level of multiplexing, and since it provides companies that of a single base extension (SBE) probe used in SNP invention in the invention of the present of multiple of companies of the method of the invention of the province is that of a single base extension (SBE) probe used in SNP invention nvention

Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;

Query Match 0.9%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 4.2e+02; Matches 24; Conservative 0; Mismatches 0; Indels 2709 AAAAAAAAAAAAAAAAAAA 2732 ò

24 AAAAAAAAAAAAAAAAAA 1 a

AAD62664 standard; DNA; 24 BP AAD62664; AAD62664/ 

12-FEB-2004 (first entry)

Immunostimulatory T-rich oligonucleotide #2183.

Antibody dependent cellular cytotoxicity; ADCC; immune response; wart; imidazoquinoline agent; asthma; allergy; infectious disease; cancer; cytostatic; antimicrobial; dermatological; virucide; ss.

Unidentified

US2003139364-A1

24-JUL-2003.

15-OCT-2002; 2002US-00272502

12-OCT-2001; 2001US-0329208P

(IOWA ) UNIV IOWA RES FOUND.

Bauer S; Krieg AM, Schetter C, Bratzler RL, Vollmer J, Jurk M,

WPI; 2003-829705/77.

Stimulating antibody dependent cellular cytotoxicity, modulating immune response and inducing antigen-specific immune response in subject by administering imidazoguinoline agents in conjunction with other agents.

Disclosure; Page 11; Opp; English.

The invention relates to methods for stimulating antibody dependent cellular cytotoxicity (ADCC), for modulating immune response and for inducing antigen-specific immune response which involve administering imidazoquinoline agents in conjunction with other agents. The method is useful for stimulating ADCC in a subject having a disorder chosen from asthma/alleryy, infectious disease, cancer and warts. The present sequence is an immunostimulatory oligonucleotide used to illustrate the method of the invention 

Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;

Gaps ö Score 24; DB 1; Length 24; Pred. No. 4.2e+02; 0; Indels 100.0%; Preα. ... 0.9%; 24; Conservative Query Match Best Local Similarity Matches

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2709 AAAAAAAAAAAAAAAAAAA 2732 24 AAAAAAAAAAAAAAAAAAAAAAA 1 g

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RESULT 282 ACA58802/c

ACAS8802 standard; DNA; 24

ACA58802;

(first entry) 10-JUN-2003 Gastric ulcer treatment immunostimulatory nucleic acid #148.

Gastric ulcer, ss; immunostimulant; equine gastric ulcer syndrome; EGUS; Heliobacter pylori

Synthetic.

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Gaps

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US2002198165-A1

26-DEC-2002.

01-AUG-2001; 2001US-00920313

01-AUG-2000; 2000US-0222248P

BRATZLER R PETERSEN D BRAT/) (PETE/)

Bratzler RL, Petersen DM;

WPI; 2003-370798/35.

Prevention or treatment of gastric ulcer involves administering nucleic 

Disclosure; Page 14; 45pp; English.

ulcer comprising administering a nucleic acid to a subject in need for treatment of gastric ulcer. A nucleic acid sample comprising oligonucleotide 2006 was administered to a mouse model by an oral route or a vehicle control. Colonisation of mice by Heliobacter pylori was assessed at time points from 1 day to 1 month after treatment. The assessed. The method is useful for preventing or treating a gastric ulcer on a subject e.g. human or non-human vertebrate animal including dog, The invention relates to a method of prevention or treatment of gastric

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cat, horse (equine gastric ulcer syndrome, EGUS), cow, goat, sheep, pig, rabbit, turkey, chicken, primate, rat and mouse. The method effectively treats or prevents gastric ulcers. The present sequence represents an immunostimulatory nucleic acid for the treatment of gastric ulcers
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Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;

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ö 0.9%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 4.2e+02; Indels 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAAAAAAA 2732 Conservative Query Match Best Local Similarity Thes 24; Conserva 24 ò 엄

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Gaps

Non-CpG DNA oligonucleotide IMT 053 SeqID 19. ADG75917 standard; DNA; 24 (first entry) 11-MAR-2004 ADG75917; RESULT 283 ADG75917, 

ss, CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytold dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.

Synthetic.

WO2003101375-A2

11-DEC-2003.

30-MAY-2003; 2003WO-EP005691.

30-MAY-2002; 2002CA-02388049.

(IMMU-) IMMUNOTECH SA

Lopez RA;

WPI; 2004-053333/05.

New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. nelanoma

Example 3; SEQ ID NO 19; 139pp; English.

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune artibody production in Brealls, as well as cell differentiation, cytokine antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacycoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an invention describes immunomodulator compositions that also comprise an enterpy for inducing B-cell activation, treating, preventing or therapy for inducing B-cell activation, treating, preventing or mellorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is the non-CpG DNA oligo IMT the invention. used in an exemplification of

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Gaps

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0; Indels

100.0%; Pred. w. 0; Mismatches

24; Conservative

Query Match Best Local Similarity

Best Loc Matches

2709 AAAAAAAAAAAAAAAAAAAAA 2732

AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

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Score 24; DB 1; Length 24; Pred. No. 4.2e+02;

Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;

ö The invention relates to a method of detection of deposition unit misalignment of an in situ polymeric array synthesis device which comprises synthesising test probe feature(s) on substrate using in situ polymeric array synthesis device, contacting test probe feature with at least two different distinguishably labelled targets and evaluating binding of labelled targets to test probe feature to detect any pulse jet misalignment of polymeric array synthesis device. The method is useful for detecting deposition unit misalignment e.g. printhead misalignment, of an in situ polymeric, e.g. nucleic acid, array synthesis device. The method is easy to use, cost effective effection and/or adjustments of misalignments and may enable immediate detection and/or adjustments of one or more printheads of an in situ nucleic acid array synthesis fluid Gaps Detection of deposition unit misalignment of in situ polymeric array synthesis device, by contacting test probe feature with different distinguishably labeled targets, and evaluating binding of labeled deposition device if misalignment is detected. The present sequence ô ss; deposition unit misalignment; polymeric array synthesis; pulse jet misalignment; printhead misalignment; microarray. represents an oligonucleotide synthesised on a microarray Score 24; DB 1; Length 24; Pred. No. 4.2e+02; 0; Mismatches 0; Indels Microarray synthesised oligonucleotide #10. 2709 AAAAAAAAAAAAAAAAAAAA 2732 Kronick MN; 24 AAAAAAAAAAAAAAAAAAAAAAAAA 1 Example 2; Page 16; 36pp; English. 0.50, 100.0%; Pretargets to test probe feature. BP 25-FEB-2003; 2003US-00374307 25-FEB-2003; 2003US-00374307 24 (first entry) DA, Query Match 0.9 Best Local Similarity 100. Matches 24; Conservative ADR48246 standard; DNA; Amorese (LEPR/) LEPROUST E M. (AMOR/) AMORESE D A. (KRON/) KRONICK M N. WPI; 2004-634540/61. US2004170984-A1. Leproust EM, 18-NOV-2004 02-SEP-2004. Synthetic. ADR48246; RESULT 284 ADR48246 a ò

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Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of detection of deposition unit
misalignment of an in situ polymeric array synthesis device which
comprises synthesising test probe feature(s) on substrate using in situ
polymeric array synthesis device, contacting test probe feature with at
least two different distinguishably labelled targets and evaluating
contacting of labelled targets for test probe feature to detect any pulse jet
misalignment of polymeric array synthesis device. The method is useful
for detecting deposition unit misalignment e.g. printhead misalignment,
of an in situ polymeric, e.g. nucleic acid, array synthesis device. The
method is easy to use, cost effective effective at detecting printhead
misalignments and may enable immediate detection and/or adjustments of
cone or more printheads of an in situ nucleic acid array synthesis fluid
deposition device if misalignment is detected. The present sequence
represents an oligonucleotide synthesised on a microarray.
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                                                                                                                                                                                                                                                                                                                                                      Detection of deposition unit misalignment of in situ polymeric array synthesis device, by contacting test probe feature with different distinguishably labeled targets, and evaluating binding of labeled targets to test probe feature.
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                                                                                                            ss; deposition unit misalignment; polymeric array synthesis; pulse jet misalignment; printhead misalignment; microarray.
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                                                                                       Microarray synthesised oligonucleotide #13
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                                                                                                                                                                                                                                                                                                           Kronick MN;
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                     ADR48249 standard; DNA; 24 BP.
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                                                                                                                                                                                                                                       25-PEB-2003; 2003US-00374307.
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                                                                  (first entry)
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                                                                                                                                                                                                                                                             (LEPR/) LEPROUST E M.
(AMOR/) AMORESE D A.
(KRON/) KRONICK M N.
                                                                                                                                                                                                                                                                                                                                WPI; 2004-634540/61.
                                                                                                                                                                    US2004170984-A1,
                                                                  18-NOV-2004
                                                                                                                                                                                          02-SEP-2004
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                                                                                                                                              Synthetic.
                                           ADR48249;
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RESULT 285
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           ADR48249/
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The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including thintis, urticaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
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ss; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibaccerial; virucide; immunoglobulin B antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.
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Pred. No. 4.2e+02;
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100.0%; Pred. No. ...
... 0; Mismatches
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02-FEB-2001; 2001US-00776479.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BRAT/) BRATZLER F
(PETE/) PETERSEN I
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                US2004235774-A1
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Petersen DM,

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Bratzler RL,
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                                                                                                                                                                                      Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     so; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Allergic response suppressor oligonucleotide #433.
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Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                               , Match 0.9%; Score 24; DB Local Similarity 100.0%; Pred. No. 4.2 nes 24; Conservative 0; Mismatches
                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 961; 235pp; English
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02-FEB-2001; 2001US-00776479.
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                         23-APR-2004; 2004US-00831778
                                                  03-FEB-2000; 2000US-0179991P.
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                                                                                                                                            Bratzler RL, Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BRATZLER R L.
PETERSEN D M.
FOURON Y.
                                                                                        BRATZLER R L.
PETERSEN D M.
                                                                                                                                                                  WPI; 2004-833006/82.
                                                                                                                 (FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004235774-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BRAT/) E
(PETE/) I
(FOUR/) I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADU89749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                         (BRAT/)
                                                                                                     PETE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 288
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The invention relates to an oligonuclectide molecule (ON) labeled with several fluorophores of one or more types embedded in its backbone, where one or more of the fluorophores is not located at either the 3' or 5' terminus of ON. ON is useful for sequencing nucleic molecules. ON is useful for detecting biological molecules e.g., antibody, antigen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                         The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, uriticaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. This sequence represents an oligonucleotide used in the
                                 Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel oligonucleotide molecule labeled with several fluorophores, useful for detecting biological molecules e.g., antibody, antigen, avidin or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fluorophore, detection; antibody; antigen; avidin; hormone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fluorophore-labeled biological detection oligonucleotide #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 4.2e+02;
rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                         Disclosure; SEQ ID NO 433; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID NO 5; 18pp; English.
                                                                               of an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24 AAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MONS ) MONSANTO TECHNOLOGY LLC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-MAY-2001; 2001US-00859736.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-MAY-2000; 2000US-0205452P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADV86472 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                    method of the invention.
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WPI; 2004-833006/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Li WR, Zhou JS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6838244-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADV86472;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amplifying RNA from a subject, useful for expression analysis, by converting RNA into circularized cDNA suitable for amplification by rolling circle replication, and transcribing the amplified CDNA into RNA, resulting in amplified RNA.
            18
avidin, protein, peptide, bacteria, virus, blood cell or hormone. ON capable of providing strong fluorescence signals at different wavelengths. This sequence corresponds to an example of an
                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents a primer which was used in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88; primer; PCR; rolling circle replication; expression analysis;
DNA replication.
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0
                                                                                                                                                                                                                                                                                     Length 24;
                                                                                                                                                                                                                                                                        Query Match 0.9%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 4.2e+02; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                   Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAAAAAAA 2732
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                                                                                                                      oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rolling replication primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-JUL-2004; 2004WO-US022997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADW99566 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Liang H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2005-152234/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADW99566;
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The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating nonallergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
                                                                                                                                                                                                                                  Immunostimulant, Antiinflammatory, Antipsoriatic; Gastrointestinal-Gen., Antiulcer; Dermatological, Antiallergic; helper T-lymphocyte; immune stimulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helper1 cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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Pred. No. 4.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "Phosphorothioate backbone"
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                                                                                                                                                                                                      Immunostimulatory oligonucleotide, SEQ ID 415.
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100.0%; Pred. No. 4...
                                                                                                                                                                                                                                                                                                  contact dermatitis; ss; phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 415; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            contact dermatitis or latex dermatitis.
2709 AAAAAAAAAAAAAAAAAAA 2732
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                  29-MAR-2001; 2001US-0279642P. 29-MAR-2002; 2002US-00112653.
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                                                                                                         AED75279 standard; DNA; 24
                                                                                                                                                                         (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                               modified base
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                                                                                                                                        AED75279;
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RESULT 292

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0.9%; Score 24; DB 1; Length 24; 100.0%; Pred. No. 4.2e+02; ative 0; Mismatches 0; Indels

Best Local Similarity 100. Matches 24; Conservative

Query Match

AED75711

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/*tag=
                               US2005250726-A1.
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         Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (1) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating non-allergic inflammatory diseases and ulcerative colitis), eczema, allergic contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; immune stimulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy;
                                                                                      Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiinfeer; Dermatological; Antiallergic; helper T-lymphocyte; immune stimulation; inflammation; psoriasis; inflammatory bowel disease; crohns disease; ulcerative colitis; eczema; skin allergy;
                                                                                                                                                                                                                                                                                                                                                Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helper1 cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunostimulatory oligonucleotide, SEQ ID 919.
                                                                  Immunostimulatory oligonucleotide, SEQ ID 920
                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 920; 58pp; English.
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29-MAR-2002; 2002US-00112653.
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                                                                                                                                   contact dermatitis; ss.
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                                                                                                                                                                                                                                                                                                                             WPI; 2005-768014/78.
                                                                                                                                                                                                                                                                                                         Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                             12-JAN-2006
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                                                                                                                                                          Synthetic.
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The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation, and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating nonallergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
                                                                                                                                                                                                                                                                                                                 Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   '*tag= a
'ote= "labelled with the donor carboxyfluoscein"
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100.0%; Pred. No. 4.2e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing primer used to exemplify the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 919; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2709 AAAAAAAAAAAAAAAAAAA 2732
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12-MAY-2005; 2005US-00127654.
                                                          29-MAR-2001; 2001US-0279642P
                                                                                   29-MAR-2002; 2002US-00112653
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Best Local Similarity 100...
Best Local Similarity 100...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag≃
                                                                                                                                             (IOWA ) UNIV IOWA RES
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treatment
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                                         Sakaki Y;
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                                                                                                                                                                                                                                                                                                                 AAX84260;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                       PCR
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                                                                                                                                                                                                                                                       The present sequence exxemplified the primer of the invention, and is used to sequence Incyte clone 1 (AAV42737). The primer of the invention is labelled with a set of at least 2 different fluorescent labels. The set comprises an energy-transfer fluorescent label with at least 1 each of a donor fluorophore and an acceptor fluorophore capable of energy transfer, and separated by a distance x, and a second similar fluorescent label in which the separation distance is y, x and y being sufficiently different for the two fluorescent labels to produce distinct fluorescent signals. Fluorescent labels are useful in multicomponent analyses, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       associated protein 1; Napl; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                Set of energy-transfer fluorescent labels with donor and acceptor at different separations - useful for DNA sequencing allows use of fewer analysing wavelengths or an increased throughput.
                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                           /*tag= b
/note= "optionally laballed with the acceptor
carboxyrhodamine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer for human Nck associated protein 1 coding sequence.
/note= "optionally labelled with the donor carboxyfluoscein"
                                                                                                                                                                                                                                                                                                                                                                                        0.9%; Score 24; DB 1; Length 25;
100.0%; Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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                                                                                                                       97US-00784162
                                                                                                    97WO-US022914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97JP-00363183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX84258 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                         (INCY-) INCYTE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.
108 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy; PCR primer; ss.
                                                                                                                                                                             WPI; 1998-414127/35.
                                                                                                                                                                                                                                                                                                                                                     for DNA sequencing
                   modified base
                                                                                                    12-DEC-1997;
                                                                                                                       15-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                   23-JUL-1998
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This sequence represents a PCR primer used to isolate DNA encoding the human Nck associated protein 1 (Napl) of the invention. Napl inhibits apoptosis. The protein can be used in the investigation, diagnosis and
                                                                                                                                                                                               Protein inhibiting apoptosis, useful in the diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nck associated protein 1; Napl; human; apoptosis; Alzheimer's disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein inhibiting apoptosis, useful in the diagnosis and treatment of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer for human Nck associated protein 1 coding sequence.
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Pred. No. 4.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                        (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 25 BP; 0 A; 0 C; 1 G; 24 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                             Example 1; Page 76; 90pp; Japanese.
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(KYOW ) KYOWA HAKKO KOGYO KK.
(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-JP005646.
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(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.98;
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                                                                                                                                                                                                                            Alzheimer's disease.
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                                                                                                                                        WPI; 1999-395181/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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Murine DNA oligonucleotide #10

11-AUG-2005

AEA31163;

AEA31163 standard; DNA; 25

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New calix(4) arene-nucleoside hybrid useful in gene therapy has at least one nucleoside attached to a calix(4) arene group through amide bonding, and is derived from a calix(4) arene having amino groups.
                            Gaps
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                            Indels
                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
/note= "OTHER= calix(4)arene nucleoside"
                                                                                                                                                                                                                                                                          Calix(4) arene; triplex; gene therapy; DNA sensor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 25 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 1 Other;
           Pred. No. 4.3e+02;
; Mismatches 0;
                                                        2709 AAAAAAAAAAAAAAAAAAAAAA 2732
                                                                                                                                                                                                                                            Calix(a) arene-oligonucleotide hybrid
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/*tag= a
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100.08; PL
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ID ACF79235 Btandard; DNA; 25 BP.
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                            24; Conservative
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Best Local Similarity
              Best Local Similarity
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                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                stem loop
                                                                                                                                                                                       ACF79235;
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Matches
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The invention relates to a synthetic mammalian (retro)transposon ORF2 or ORF1 gene exhibiting a higher level of expression relative to a natural CC ORF1 gene. The invention also relates to a (retro)transposon Comprising the synthetic gene, a mammalian L1 crerotransposon comprising the synthetic gene, a recombinant vector construct comprising the synthetic gene, a eukaryotic cell transfected, cransformed or infected with the recombinant vector construct comprising the synthetic gene, a eukaryotic cell transfected, construct comprising the synthetic gene, a eukaryotic cell transfected, confidentifying a dealired gene, or its biologically active fragment, to the cells of a mammal, a composition comprising a cassette comprising the composition is useful for treating a genetic disorder in a cells. The composition is useful for treating a genetic disorder in a mammal such as hemophilia, Parkinsons disease, Pabry disease, confidentifying an uncharacterized gene, or its biologically active fragment, confidently disorders disease, cystic fibrosis, disease, denosine deaminase deficiency, alpha-antitrypsin confidency, buchenne muscular dystrophy, phenylectonuria, sickle cell diseases. This sequence represents a murine DNA oligonucleotide used in
                                                                                                                                                                                                                                       Tay Sachs disease; thalassemia; lysosomal storage disease; metabolic disorder; antiparkinsonian; hemostatic; metabolic; antilipemic; CNS-Gen.; respiratory-gen.; antianemic; cerebroprotective; muscular-gen.; dermatological; nootropic; antisickling; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New synthetic mammalian (retro)transposon open reading frame 2 (ORF2) or ORF1 gene exhibiting a higher level of expression relative to a natural 11 (retro)transposon ORF2 or ORF1 gene, useful for treating e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                  Parkinsons disease; Fabry disease; hypercholesterolemia; Gauchers disease; cystic fibrosis; adrenolemcoystrophy; adenosine deaminase deficiency; alpha-1 antitrypsin deficiency; buchenne dystrophy; phenylketonuria; sickle cell anemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                              Transposon; retrotransposon; genetic disorder; hemophilia;
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100.0%; Pred. No. 4.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                   WO2005049789-A2.
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RESULT 298

**AEA31163** 

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The invention relates to a synthetic mammalian (retro)transposon ORF2 or ORF1 gene exhibiting a higher level of expression relative to a natural inferro)transposon ORF2 or ORF1 gene. The invention also relates to a cretro transposon comprising the synthetic gene, a mammalian Liretro)transposon comprising the synthetic gene, a mammalian Liretrotransposon comprising the synthetic gene, a recombinant vector construct comprising the synthetic gene, a ewkaryotic cell transfected, transformed or infected with the recombinant vector construct, a method of delivering a desired gene, or its biologically active fragment, to the cells of a mammal, a composition comprising a cassette comprising the gene, desired gene and a pharmaceutical carrier, and a method of identifying an uncharacterized gene, or its biologically active fragment, in cells. The composition is useful for treating a genetic disorder in a mammal such as hemophilia, Parkinsons disease, Fabry disease, hypercholesterolemia, Gauchers disease, cystic fibrosis, adrenoleukodystrophy, disorders associated with mutations in the dystrophin gene, adenosine deaminase deficiency, alpha-antitrypsin anemia, Tay Sachs disease, thalassemia, lysosomal storage diseases and metabolic disorders. The synthetic gene is useful for treating the charactering an emaia, Tay Sachs disease, thalassemia, lysosomal storage diseases and the contraction of the charactering the characterin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tay Sachs disease; thalassemia; lysosomal storage disease; metabolic disorder; antilipemic; metabolic disorder; antilipemic; CNS-Gen.; respiratory-gen.; antilanemic; cerebroprotective; muscular-gen.; dermatological; nootropic; antisickling; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New synthetic mammalian (retro)transposon open reading frame 2 (ORF2) or ORF1 gene exhibiting a higher level of expression relative to a natural L1 (retro)transposon ORF2 or ORF1 gene, useful for treating e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transposon; retrotransposon; genetic disorder; hemophilia; Parkinsons disease; Fabry disease; hypercholesterolemia; Gauchers disease; cystic fibrosis; adrenoleukodystrophy; adenosine deaminase deficiency; alpha-1 antitrypsin deficiency; Duchenne dystrophy; phenylketonuria; sickle cell anemia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; SEQ ID NO 15; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murine DNA oligonucleotide #11.
                                                                                                                AEA31164 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAY-2004; 2004WO-US015810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAY-2003; 2003US-0473658P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                metabolic diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2005049789-A2.
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                                                                                                                                                                                                                                      AEA31164;
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RESULT 299
                                                                    AEA31164
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The invention relates to a synthetic mammalian (retro) transposon ORF2 or ORF1 gene. The invention also relates to a natural L1 (retro) transposon ORF2 or ORF1 gene. The invention also relates to a (retro) transposon comprising the synthetic gene, a mammalian L1 construct comprising the synthetic gene, a recombinant vector or tetrotransposon comprising the synthetic gene, a recombinant vector construct comprising the synthetic gene, a recombinant vector construct comprising the synthetic gene, a casentrock a method of delivering a desired gene, or its biologically active fragment, to the cells of a mammal, a composition comprising a cassette comprising the gene, a desired gene and a pharmaceutical carrier, and a method or identifying an uncharacterized gene, or its biologically active fragment, in cells. The composition is useful for treating a genetic disorder in a mammal such as hemophilia, Parkinsons disease, Fabry disease, hypercholesterolemia, Gauchers disease, rebry disease, cystic fibrosis, adrenoleukodystrophy, disorders associated with mutations in the deficiency, buchenne muscular dystrophy, phenylectonuria, sickle cell metabolic disporders. The synthetic gene is useful for treating the chases and metabolic disporacers. The synthetic gene is useful for treating the diseases. This sequence represents a murine DNA oligonucleotide used in the cope of the invention.
                                                                                                                                                                                                                                                                               Transposon; retrotransposon; genetic disorder; hemophilia; Parkinsons disease; Fabry disease; hypercholesterolemia; Gauchers disease; cystic fibrosis; adrenoleukodystrophy; adenosine deaminase deficiency; alpha-1 antitrypsin deficiency; Duchenne dystrophy; phenylketonuria; sickle cell anemia; Tay Sachs disease; thalassemia; lysosomal storage disease; metabolic disorder; antiparkinsonlan; hemostatic; metabolic; antilipemic; CNS-Gen.; respiratory-gen.; antilianemic; cerebroprotective; muscular-gen.; dermatological; nootropic; antisickling; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New synthetic mammalian (retro)transposon open reading frame 2 (ORF2) or ORF1 gene exhibiting a higher level of expression relative to a natural Li (retro)transposon ORF2 or ORF1 gene, useful for treating e.g., metabolic diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 13; 66pp; English.
1 AAAAAAAAAAAAAAAAAAA 24
                                                                                                                                                                                                                                        Murine DNA oligonucleotide #9.
                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-MAY-2004; 2004WO-US015810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAY-2003; 2003US-0473658P.
                                                                                                       AEA31162 standard; DNA; 25
                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Han JS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2005049789-A2
                                                                                                                                                                                                11-AUG-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUN-2005.
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                                                                                                                                                  AEA31162;
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Score 24; DB 1; Length 25; Pred. No. 4.3e+02;

100.0%;

Query Match Best Local Similarity

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Gaps

; 0

0.9%; Score 24; DB 1; Length 25; 100.0%; Pred. No. 4.3e+02; ative 0; Mismatches 0; Indels

Query Match
Best Local Similarity 100.0
Matches 24; Conservative

Sequence 25 BP; 24 A; 0 C; 0 G; 1 T; 0 U; 0 Other;

Matches

RESULT 301

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This invention describes a novel method (M1) for inserting a nucleic acid (N1) into a circular vector (V1) comprising joining ends of N1 and V1 under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reannealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex populations of DNA or cDNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger nucleic acid fragments are just as easily cloned, allowing highly representative libraries to be made. Vector to vector ligation is avoided using the methods. AAA40351-A40366 represents primers used to illustrate
                                                                                                                                                                                                                                                                                                                                         Inserting a nucleic acid into a circular vector comprising joining their ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 24; DB 1; Length 28;
Pred. No. 4.5e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 28 BP; 1 A; 1 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pBluescriptSK+ phagemid primer SEQ ID NO: 12.
 Θ.
pBluescriptSK+ phagemid primer SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 67; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.
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                                     Primer; cloning; ligation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US030277.
                                                                                                                                                                         99WO-US030277.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer; cloning; ligation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-NOV-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 100.
nes 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA40362 standard; DNA;
                                                                                                                                                                                                                                           (ROMA/) ROMANTCHIKOV
                                                                                                                                                                                                                                                                                                           WPI; 2000-442381/38.
                                                                                                                                                                                                                                                                              Romantchikov Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200036088-A1
                                                                                                       WO200036088-A1
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                                                                                                                                                                         17-DEC-1999;
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                                                                                                                                         22-JUN-2000
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                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                  libraries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotides comprising cholane-3,24-diol(3 alpha, 5 beta) unit which are easily absorbed into the cells, have stable structure, and form hairpin loop structure useful in RNA interference or antisense application.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to novel oligonucleotides comprising cholane-3,24-diol(3alpha,5beta) unit are provided, which are easily absorbed into cells, have a stable structure, and form a hair-pin loop structure, so that they can be used for antisense/antigene therapy or RNAi (RNA interference). The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
     Gaps
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       Indels
                                                                                                                                                                                                                                                                                                ss; RNA interference; antisense therapy; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 1 Other;
     ;
                                                                                                                                                                                                                                                                                                                                                                                                                                      'note = "OTHER=cholane-3,24-diol"
                                                                                                                                                                                                                                                              Oligonucleotide of the invention SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
       Mismatches
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                                       2709 AAAAAAAAAAAAAAAAAAAA 2732
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                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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/mod base= OTHER
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                                                                                                                                                            BP.
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                                                                                                                                              1371/c
AEC63371 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kim BH, Kim SJ;
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                                                                                                                                                                                                                              (first entry)
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       24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                    Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24;
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                                                                                                                                                                                                                                                                                                                                  Synthetic.
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RESULT 302 AAA40358/c

Matches

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Gaps

Detecting an analyte in a sample comprises providing nucleic acid sequence which is catalytically active in presence of analyte, contacting catalytic nucleic acid with substrate and amplifying catalytic product.

The invention relates to a method of detecting an analyte in a sample

Disclosure; Page; 36pp; English.

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Inserting a nucleic acid into a circular vector comprising joining their ends, melting, and reannealing ends at two different concentrations,
                                               ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
                                                                                                                                                                                                                                                                          ВР.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98IL-00126731.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    99WO-IL000557
                                                                                                                                                                                                                                                                         AAA57856 standard; DNA; 28
                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
24. .28
                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (INTE-) INTELLIGENE LTD
(ROMA/) ROMANTCHIKOV Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wathan A, Ellington A;
                           WPI; 2000-442381/38.
             Romantchikov Y;
                                                                                                                                                                                                                                                                                                                                                                                                                        WO200024931-A2
                                                                                                                                                                                            Query Match
Best Local Simi:
Matches 24;
                                                                                                                                                                                                                                                                                                     11-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                             misc_binding
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                                                              libraries
                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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misc_RNA
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This invention describes a novel method (MI) for inserting a nucleic acid (NI) into a circular vector (VI) comprising joining ends of NI and VI under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reannealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex populations of DNA or cDNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger nucleic acid fragments are just as easily cloned, allowing highly uspresentative libraries to be made. Vector to vector ligation is avoided the methods. AAA40331-A440366 represents primers used to illustrate Gaps ö Similarity 100.0%; Score 24; DB 1; Length 28; Similarity 100.0%; Pred. No. 4.5e+02; Conservative 0; Mismatches 0; Indels Sequence 28 BP; 0 A; 2 C; 2 G; 24 T; 0 U; 0 Other; 2709 AAAAAAAAAAAAAAAAAA 2732 Example 4; Page 68; 71pp; English

The method comprises providing an object of sequence which is initially catalytically inactive, but which becomes catalytically active in the presence of an analyte (the effector); providing a nucleic acid substrate for the catalytic activity of the nucleic acid sequence; and contacting the nucleic acid sequence and the substrate with the sample under conditions allowing catalytic activity of nucleic acid sequences. The conditions allowing catalytic activity of nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid sequence will be able to convert the nucleic acid product only if the analyte of interest is present. The nucleic acid acadalytic product indicates the presence of an analyte in a sample in diagnostic cassays. The invention describes the in vitro selection of a ribozyme ligase (II; AAA57859, AAA57860) which is catalytically active only in the presence of an oligonucleotide effector (AAA57854). The Li ribozyme ligase was selected from a pool of RNA molecules comprising a central constant sequence regions (the NBO RNA AB57851). In the presence of the activity of the activi

28 AAAAAAAAAAAAAAAAA S

of the effector, selection was performed using one of the targets substrate molecules AAA57855-A57857. RNAs with ligase activity (i.e., those which have become ligated to the substrate molecule) were reverse transcribed using the effector oligo, and then PCR amplified using the effector and a DNA primer identical in sequence to the substrate used for the selection. A ribozyme ligase, ii, was selected via this procedure. Ligan only adopt its active conformation (AAA57859) in the presence of the effector oligo (analyte). In the absence of the effector, ii adopts an inactive conformation (AAA57860). The present sequence represents the decoxy-T22-tagged substrate oligonucleotide. The dT22 tag enables successfully ligated products to be isolated using oligo(AA) cellulose Type 7. Note: The present sequence is not given in the specification, but

created from the inforamtion given on page 11

/bound\_moiety= "Bases 13-17 of N90 RNA pool (AAA57851)" Ribozyme; catalytic RNA; analyte detection; effector molecule; nucleic acid substrate; in vitro selection; ribozyme ligase; conformation dependent activity; allosteric activation; ss. Deoxy-T22-tagged substrate oligonucleotide. Location/Qualifiers

WPI; 2000-350763/30.

ö Cancer; cytostatic; gene therapy; triplex forming oligonucleotide; ds. Gaps ö 0.9%; Score 23.8; DB 1; Length 28; 12.6%; Pred. No. 4.7e+02; 2; Indels Sequence 28 BP; 1 A; 2 C; 1 G; 22 T; 2 U; 0 Other; 0; Mismatches 2704 GTACTAAAAAAAAAAAAAAAAAA 2730 Triplex forming oligonucleotide #4. AAL44903 standard; DNA; 29 BP. 03-MAR-2000; 2000KR-00010744. 03-MAR-2000; 2000KR-00010744 92.68; (first entry) 25; Conservative Best Local Similarity KR2001086830-A. 05-AUG-2002 Unidentified 15-SEP-2001. 27 AAL44903; Query Match 305 RESULT 305 AAL44903/c Matches à 셤 

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AAV59215
ID AAV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents an oligonucleotide used in an example of the present invention. The present invention describes a method for synthesising a selected oligonucleotide (1) having well defined ends. The method comprises: (a) annealing a primer to a single-stranded (ss) circular template to yield a primed circular template, where the template comprises: (i) at least one nucleotide sequence complementary to (1); and (ii) at least one nucleotide effective to produce a cleavage site in the oligonucleotide multimer; (b) combining the primed circular template with at least two types of nucleotide triphosphates and a polymerase enzyme with the addition of auxiliary proteins to yield a ss oligonucleotide multimer complementary to the circular oligonucleotide template, comprising multiple copies of (1); and (c) cleaving the oligonucleotide
                                                                                                                                                                                    The present invention relates to a triplex forming oligonucleotide which specifically binds to a specific gene. This is useful for the gene therapy of cancer by binding itself to Auger electron emitters. The present sequence is a triplex forming oligonucleotide of the invention
                                                                                                    Novel triplex forming synthetic oligonucleotide, useful for gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rolling circle synthesis of oligo:nucleotide(s) - using primed circular template to produce oligonucleotide multimer for cleavage.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthesis, selection, amplification, circular oligonucleotide, rolling circle synthesis, diagnosis, therapeutic agent, 88.
                                                                                                                                                                                                                                                                                                      0.9%; Score 23.8; DB 1; Length 29; 92.6%; Pred. No. 4.8e+02; tive 0; Mismatches 2; Indels
                                 Son JW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO:5 from US5174320 Example 2.
                                                                                                                                                                                                                                                                         Sequence 29 BP; 0 A; 1 C; 3 G; 25 T; 0 U; 0 Other;
                               Park MG,
                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2735
(KOCH-) KOREA CHUNGANG EDUCATIONAL FOUND
                                                                                                                                                                                                                                                                                                                                                                                                  Park GH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Col 45; 38pp; English.
                                                                                                                                                      Claim 4; Page 11; 13pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93US-00047860.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00393439
                                 ĠΧ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV12482 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYRP ) UNIV ROCHESTER.
                                 Lee DH, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-144278/13.
                                                                   WPI; 2002-233771/29.
                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5714320-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV12482;
                                                                                                                                                                                                                                                                                                             Query Match
                                 Choi JG,
                                                                                                                        tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kool ET;
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The circular template was used for the synthesis of the oligomer dT12 in an example of the method of the invention for synthesising an RNA oligonucleotide, comprising combining a single-stranded circular oligonucleotide template comprising at least one copy of a nucleotide sequence complementary to the sequence of the desired RNA oligonucleotide with at least 2 types of ribonucleotide triphosphate and a polymerase enzyme to yield a single-stranded RNA oligonucleotide multimer comprementary to the circular oligonucleotide template, where the RNA oligonucleotide multimer comprises multiple copies of the desired RNA oligonucleotide. The methods can be used for producing RNA oligonucleotides having a specific sequence and well defined ends. The RNA oligonucleotides produced can be used as probes, standards and
                                                                                                                                                                                                                                           ö
multimer at the cleavage site to produce (I) having well defined ends. The method is used for the large-scale synthesis of DNA and RNA oligomers for use, e.g. as probes and diagnostic agents and/or therapeutic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= b
/bound_moiety= "Position 26 optionally bound to position
1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
/bound_moiety= "Position 1 optionally bound to position
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; circular; cyclic; RNA oligonucleotide; probe; standard; diagnostic;
                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthesis of oligo:nucleotide(s) - using a single-stranded circular oligo:nucleotide template ribonucleotide tri:phosphate(s) and a polymerase to form multimer(s) which can be cleaved.
                                                                                                                                                                                                                                              ö
                                                                                                                                                                                    Length 26;
                                                                                                                                                                                                                                           1; Indels
                                                                                                                          Sequence 26 BP; 24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                 0.9%; Score 23.4; DB 1;
96.0%; Pred. No. 4.8e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Circular template for linear oligomer dT12.
                                                                                                                                                                                                                                                                                                     2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                               2 AAAAAAAAAACAAAAAAAAA 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 36; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-00805631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98WO-US003784.
                                                                                                                                                                              . 86.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV59215 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                  Best Local Similarity 96.0
Matches 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYRP ) UNIV ROCHESTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapeutic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9838300-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-OCT-2004
14-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV59215;
                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kool ET;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 307
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2 AAAAAAAAAAAAAAAAAAA 26

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RESULT 309

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The present invention describes a detectably labelled oligonucleotide multimer, comprising multiple contiguous copies of a repeated oligonucleotides. The detectably labelled oligonucleotide multimer is useful for detecting a target molecule. Oligonucleotide multimers may be produced in sufficient quantity to be useful for diagnostic and medical applications. The multimers are useful for affinity labelling of proteins, and for signal amplification in highly sensitive affinity capture and sequence identification in highly sensitive affinity faster, cheaper and simpler way for large-scale production of DNA and RNA oligomers and multimers. The incorporation of labels enables the oligonucleotide multimers to be useful in diagnostics and medicine. The
                                                                   Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New detectably labelled oligonucleotide multimer, comprising multiple contiguous copies of a repeated oligonucleotide - useful for detecting target molecules in diagnosis and medicinal applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present sequence represents an oligonucleotide used in an example from the present invention
diagnostic or therapeutic agents. They can be used for modifying the structure or function of a target molecule. They can also be used to cleave disease-associated RNA, DNA or protein
                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Multimer; probe; diagnosis; synthesis; detection; polymerase; ss.
                                                                                                                                                                               ö
                                                                                                                                          Length 26;
                                                                                                                                                                             1; Indels
                                                                                                      Sequence 26 BP; 24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 26 BP; 24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                       0.9%; Score 23.4; DB 1;
96.0%; Pred. No. 4.8e+02;
iive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Precircle DNA oligonucleotide SEQ ID NO:5.
                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                  Example 2; Page 41; 103pp; English.
                                                                                                                                                                                                                                                                                                                                   AAX30018 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US016776.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-00910632.
                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                           24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYRP ) UNIV ROCHESTER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-181062/15.
                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9909216-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                             ~
                                                                                                                                                                                                                                                                                                                                                                       AAX30018;
                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kool ET;
                                                                                                                                                                         Matches
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RNA oligonucleotide synthesis; ribonucleotide triphosphate; polymerase; electropozation; calcium phosphate treatment; lipid-mediated delivery; cation-mediated delivery; bacterial infection; viral infection; drug resistant infection; double stranded DNA oligomer; ss.
                                                                                                                                                                                                                                                                                                                                      Synthesizing RNA oligonucleotide involves combining single-stranded circular oligonucleotide, ribonucleotide triphosphate and polymerase enzyme to yield desired RNA complementary to circular oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                        Example 2; SEQ ID NO 5; 78pp; English.
         ВР
                                                                                                                                                                                                                        15-APR-1993; 93US-00047860.
23-FEB-1995; 95US-00393439.
26-FEB-1997; 97US-00805631.
11-MAY-2000; 2000US-00569344.
                                                                                                                                                                                                       30-NOV-2001; 2001US-00997931,
        ADC65872 standard; DNA; 26
                                                 (first entry)
                                                                     DNA oligonucleotide #5.
                                                                                                                                                                                                                                                                           (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                    WPI; 2003-755141/71
                                                                                                                                                             US2003087241-A1
                                                 18-DEC-2003
                                                                                                                                                                                   08-MAY-2003.
                                                                                                                                           Synthetic.
                            ADC65872;
                                                                                                                                                                                                                                                                                                Kool ET;
                                                                                                                                                                                                                                                                                                                                                                        template
ADC65872
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oligonuclectide, comprising combining a single-stranded circular oligonuclectide, comprising combining a single-stranded circular oligonuclectide template with at least two types of ribonucleotide triphosphate and a polymerase enzyme to yield a single-stranded RNA oligonuclectide multimer comprises multiple template, where the RNA oligonuclectide multimer comprises multiple copies of the desired RNA oligonuclectide. The method is useful for synthesising an RNA oligonuclectide with well-defined ends. The circular oligonuclectide is introduced into the cell using direct injection, calcium phosphate treatment, lipid-mediated delivery, or cation-mediated delivery. The method is useful for treating bacterial and/or viral infections in mammals, particularly drug resistant infections, and for producing double stranded DNA oligomers. The method is useful for treating addition of auxiliary proteins. This sequence represents an addition of auxiliary proteins. This sequence represents an oligonucleotide primer, or without the oligonucleotide used in the method of the invention. The invention relates to a method for synthesising an RNA

Sequence 26 BP; 24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö ch 0.9%; Score 23.4; DB 1; Length 26; 1 Similarity 96.0%; Pred. No. 4.88+02; 24; Conservative 0; Mismatches 1: ThADDE Query Match Best Local Similarity Matches

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RESULT 310

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Gaps

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0.9%; Score 23.4; DB 1; Length 26; 96.0%; Pred. No. 4.8e+02; tive 0; Mismatches 1; Indels

2709 AAAAAAAAAAAAAAAAAAAAAA 2733

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Conservative

Query Match Best Local Similarity Thes 24; Conserva

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The invention relates to an isolated polypeptide with aminopeptidase activity and the gene encoding it (including sequences complimentary to the gene and which hybridise to it at high stringency), from Aspergillus niger. Also included are a nucleic acid construct comprising the above construct comprising the above polymucleotide operably linked to one or more control sequences that direct the production of the polypeptide in a suitable expression host, a recombinant vector comprising the above nucleic acid construct, a recombinant host cell comprising the above construct or vector, and producing the protein comprising the above construct or vector, and producing the protein comprising the above construct or vector, and producing the protein comprising the above construct or vector, and producing the protein comprising the above construct or the polypeptide and recovering the polypeptide. The aminopeptides is cheese to enhance the flavour. The invention provides a bacterial enzyme for protein chydrolysis i.e. with aminopeptidase activity, to produce flavouring agents, and the enzyme has been isolated and characterised, compared to a previously observed weak aminopeptidase activity which was detected in an Aspergillus niger culture filtrate but the source was never isolated or identified. The use of enzymes to produce flavouring agents from protein manager culture filtrate but the source was never isolated or dentified the amino acids obtained. The present sequence is a reverse transcriptase (RT)-PCR primer used to investigate the intron-exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An isolated polypeptide with aminopeptidase activity, for preparing food compositions, such as bread and cheese, with enhanced flavoring.
                                                                                                                                                                                       Aminopeptidase; primer; ss; food composition; dough; flavour enhancer; baked product; cheese; poly-T; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Visser J;
                                                                                                                                                   Aspergillus niger aminopeptidase RT-PCR primer poly-T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schaap PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 BP; 0 A; 0 C; 0 G; 23 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schuurhuizen PW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              structure of the aminopeptidase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 40; 94pp; English.
                    ABK48140 standard; DNA; 24 BP.
                                                                                                                                                                                                                                                                                                                                                                                             22-AUG-2001; 2001WO-EP009925.
                                                                                                                                                                                                                                                                                                                                                                                                                                        23-AUG-2000; 2000EP-00202995
                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Basten D, Dekker PJT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-257917/30.
                                                                                                                                                                                                                                                                                                         WO200216618-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (STAM ) DSM NV.
                                                                                                          18-JUN-2002
                                                                                                                                                                                                                                                                                                                                                    28-FEB-2002
                                                                                                                                                                                                                                                                Synthetic.
                                                                 ABK48140;
ABK48140/c
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Gaps ; 0 0.8%; Score 23.2; DB 1; Length 24; 95.8%; Pred. No. 4.7e+02; ative 1; Mismatches 0; Indels Ouery Match
Best Local Similarity 95.55
Fig. 23; Conservative

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AEB90558 standard; DNA; 25 BP. AEB90558 RESULT 311 **AEB90558/** 

Thielavia terrestris NRRL 8126 cDNA library primer SEQ ID NO

20-OCT-2005 (first entry)

degradation; cellulose; glycolysis; NRRL 8126; PCR; primer; 88; CDNA library detergent; 

Escherichia coli.

WO2005074647-A2.

18-AUG-2005.

28-JAN-2005; 2005WO-US003525.

30-JAN-2004; 2004US-0540661P

(NOVO ) NOVOZYMES INC.

Mcfarland K; Vlasenko E, Re E, Zaretsky E, Brown K, Harris P, Lopez De Leon A;

WPI; 2005-555779/56.

New polypeptide having cellulolytic enhancing activity, useful for degrading or converting biomass to sugars and for producing organic substance and detergent composition.

Example 11; SEQ ID NO 43; 219pp; English.

The invention describes an isolated polypeptide (I) having cellulolytic enhancing activity. Also described are: producing a polymucleotide having a mutant nucleotide sequence, an isolated polymucleotide comprising a complexity sequence, an isolated polymucleotide compression by the method of (1); a nucleic acid construct comprising the complexity linked to one or more control sequences that direct the production of the polypeptide in an expression host or a conclectide sequence encoding a signal peptide consisting of nucleotides 330-387 of SEQ ID NO: 1, nucleotides 47-97 of SEQ ID NO: 3, nucleotides 69-125 of SEQ ID NO: 9, where the gene is foreign to the nucleotides sequence; a recombinant expression vector comprising the nucleotide sequence; a recombinant expression vector comprising the conclection construct of (3); producing (1); producing a mutant cell; a mutant cell produced by the method of (7); producing a protein; a transgenic plant, plant part or plant cell, which has been comprising the polypeptide having cellulolytic enhancing activity, a cellulolytic activity, and a surfaceant; degrading or converting a colludorytic phancing and producing an organic cubstance. (1) is a of 125, 478, 516, 452, or 608 amino acids (EVEN SEQ ID NOS: 2-10) encoded by a polynucleotide comprising a nucleotide sequence of 1846, 880, 1000, 681, or 960 bp (0DD SEQ ID NOS: 1-9). The polypeptide, polynucleotide, and methods are useful for degrading or converting biomass to sugars, for producing an organic substance, and for producing detergent composition. This sequence represents a primer used in the creation of a Thielavia terrestris NRRL 8126 cDNA library for the isolation of cellulolysis polypeptide having cellulolytic enhancing activity comprising a sequence of 326, 478, 516, 452, or 608 amino acids (EVEN SEQ ID NOS: 2-10) encode

Sequence 25 BP; 0 A; 0 C; 0 G; 23 T; 0 U; 2 Other;

enhancing proteins

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Gaps ö ch 0.8%; Score 23.2; DB 1; Length 25; 1. Similarity 95.8%; Pred. No. 4.8e+02; 23; Conservative 1; Mismatches 0; Indels Query Match Best Local Similarity Best Loca Matches

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Immunostimulatory non-CpG phosphorothioate DNA oligo IMT191.
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                                                                                                                                                          11-DEC-2003.
                                                                                                              Synthetic.
                                                                                                                                                                                                                                                              Lopez RA;
                                                                                                                                                                                                                                                                                                                                                      melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         XEXEXEX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA coligonucleotides (without a CPG motif.) which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present in plasmacytoid dendritic cells. The present cinvention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is a non-CpG DNA oligo of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                 New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                   ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 23.2; DB 1; Length 28; 89.3%; Pred. No. 5.1e+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 28 BP; 1 A; 1 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAAAAAAAAAAAA 2736
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 17; Page 82; 139pp; English.
                                                                                             Non-CpG DNA oligonucleotide 61.
                    ADG76060 standard; DNA; 28 BP.
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                                                                                                                                                                                                                                                                                                30-MAY-2002; 2002CA-02388049.
                                                                                                                                                                                                                                                                       30-MAY-2003; 2003WO-EP005691
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                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                        (IMMU-) IMMUNOTECH SA
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-053333/05.
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Best Local Similarity
Matches 25; Conserv
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                                                                    11-MAR-2004
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                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention.
                                           ADG76060;
                                                                                                                                                                                                                                                                                                                                                  Lopez RA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         melanoma.
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This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA coligonucleotides (without a CPG motif), which can stimulate an immune coligonucleotides (without a CPG motif), which can stimulate an immune response in animals of the order of prinate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendrific cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, virtues, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or cells and glycolipids. As such, these DNA oligos can be used in gene comprised in munne system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory corrionate non-CPG variant DNA oligo, used to determine the effect of oligo size on B cell proliferation and ILG secretion in an exemplification of the invention. NOTE: This sequence is referred to as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
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ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differation; differation; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
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tes 25; Conservative
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                                                                                                                                                                                        renal cell carcinoma
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WPI; 2000-664908/64.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and assay.
                                                                                                                                                                                                                                                                                                                                              ADT55093;
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                                                                                                                                                                                                               Query Match
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ADT55093
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                                                                                                                                                                                                                                                                        요
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                                                                                                                                                                                                                       Detaching nucleic acid molecule comprising unconventional nucleotide incorporated at predetermined site from a solid support involves cleaving the nucleic acid molecule at the site of unconventional nucleotide.
                                                                                                                                                                                                                                                                                 The present invention is concerned with the cleavage of nucleic acids from solid supports. This is carried out by adding a non-conventional nucleotide into the nucleic acid attached to the support, so that it is recognised and cleaved by a specific DNA glycosylase and the sequence is released. This is useful in many molecular biological procedures such as sequencing, in vitro amplifications, CDNA and template preparation, DNA-based assays, mutagenesis procedures, nucleic acid purification and affinity chromatography. The present sequence is an oligonucleotide used in assays to demonstrate the methods of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acids from solid supports assay oligonucleotide #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
   Nucleic acid cleavage; solid support; DNA-RNA hybrid;
affinity chromatography; sequencing; mutagenesis; DNA preparation;
nucleic acid purification; ss.
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.00.0%; Pred. No. 4.7e+02;
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 acid cleavage; solid support; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23 BP; 0 A; 0 C; 0 G; 22 T; 1 U; 0 Other;
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Best Local Similarity 100.
Matches 23; Conservative
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                                                                                 /*tag=
                                                                                                                                                                                    (GOLD/) GOLDSBOROUGH
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                                           Synthetic
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                                                                        misc RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention comprises an electrophoresis apparatus for isolating variant genes. The apiarists consists of: a sealed flow path filled with buffer contealing linear polymer and DNA joint controlling agent, and containing DNA conjugate for separation, DNA conjugate for presention, DNA conjugate for present on the conjugate for assay, and a heating apparatus for heating the portion of sealed flow path in which the DNA conjugate for sasy is fixed. The electrophoresis apparatus is useful for isolating a variant gene. The electrophoresis apparatus is also useful in gene diagnosis for
Detaching nucleic acid molecule comprising unconventional nucleotide incorporated at predetermined site from a solid support involves cleaving the nucleic acid molecule at the site of unconventional nucleotide.
                                                                                                                                                                                                                                                       The present invention is concerned with the cleavage of nucleic acids from solid supports. This is carried out by adding a non-conventional nucleotide into the nucleic acid attached to the support, so that it is recognised and cleaved by a specific DNA glycosylase and the sequence is released. This is useful in many molecular biological procedures such as sequencing, in vitro amplifications, cDNA and template preparation, DNA-based assays, mutagenesis procedures, nucleic acid purification and affinity chromatography. The present sequence is an oligonucleotide used in assays to demonstrate the methods of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ch 0.8%; Score 23; DB 1; Length 23; 1 Similarity 100.0%; Pred. No. 4.7e+02; 23; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 0 A; 0 C; 0 G; 0 T; 23 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAAAAAAA 2731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                           Example 1; Page 32; 47pp; English.
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Compound activity characterisation-related oligonucleotide SegID4.

(first entry)

26-FEB-2004

ADG16129;

ADG16129 standard; DNA; 24

RESULT 318

ADG16129/

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detecting the presence or absence of a gene mutation, or variant gene, and for single nucleotide polymorphism analysis. The present DNA sequence was shown in a figure exemplifying the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Electrophoresis apparatus useful for isolating variant gene, comprises heating apparatus, and sealed flow path comprising linear polymer and DNA joint controlling agent, with DNA conjugates for separation, purification
                                                                                                                                                            Gaps
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gene mutation detection; variant gene detection;
single nucleotide polymorphism analysis; SNP detection; ss.
                                                                                                                      Length 23;
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                                                                             Sequence 23 BP; 23 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                        4.7e+02;
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larity 100.0%; Pred. No. 4.7e+02;
Conservative __0; Mismatches 0;
                                                                                                                   ch 0.8%; Score 23; DB 1;
1 Similarity 100.0%; Pred. No. 4.7e+0
23; Conservative 0; Mismatches
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                                                                                                                                    Local Similarity
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This invention relates to a novel method for the characterisation of the activity of a compound on cell. The method involves receiving images of cells with a cellular activity and images of other cells treated with the compound, quantitatively characterising phenotypic attributes of the image of cells with a cellular activity to produce a target phenotype for the cellular activity and that of the image of other cells to produce a second phenotype for the compound, and comparing the two phenotypes to determine whether the compound, and comparing the two phenotypes to determine whether the compound possesses cellular activity. The invention may be useful for characterising cellular activity of a compound, for determining information about properties of substances based upon the information about structure of living or non-living cells exposed to substances. The invention is also useful for identifying promising multiple biological descriptors from a single cell markers or components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Characterizing cellular activity of compound, by receiving images of cells with known activity and images of cells treated with compound, characterizing phenotypic attributes of images and comparing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                           Crompton AM;
                                                                                                                               compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate treatment; multiple biological descriptor; cell marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         Vaisberg EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 0 A; 1 C; 0 G; 23 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST polymorphic DNA repeat polynucleotide #134.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 23; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. No. 4.8 ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                         Adams CL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 18; 139pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                     26-APR-2000; 2000US-0199778P.
20-FEB-2001; 2001US-00790214.
                                                                                                                                                                                                                                                                                                      24-APR-2001; 2001WO-US013248.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.88;
                                                                                                                                                                                                                                                                                                                                                                                                                         Destreicher DR, Sabry JH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; cDNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                        (CYTO-) CYTOKINETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-041423/05.
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Best Local Similarity
                                                                                                                                                                                                                                 WO200181895-A2.
                                                                                                                                                                                                  Unidentified.
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                                                                                                                                                                                                                                                                   01-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phenotypes.
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Local Similarity les 23; Conserv

Best Loca Matches

Query Match

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23

Length 23; 0; Indels US2003121063-A1.

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repeat within a coding sequence (expressed sequence tag, EST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic candidate polymorphic repeat. The computational methods (polymorphic conditions) in the sequences, powpous, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River CC syndromic Haussian diseases, fragile-X syndrome, Fredreich s ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying a candidate polymorphic repeat within a coding sequence, for understanding or treating genetic disease, comprises detecting tandem repeats in a target coding sequence and scoring the repeats for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention discloses a method for identifying a candidate polymorphic
            EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPOUS; Rep-X; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy; insertional mutation; germ line specific promoter; mutation generation; transgenic animal; poly A element; non-LTR; retrotransposon; long terminal repeats; human; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        L1 retrotransposon insertion characterisation primer seg id 151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 0 A; 1 C; 0 G; 23 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 4.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                              Fondon JW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Prec. ...
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                                                                                                                                                                                                                                                                                                                                                                                                              Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF12405 standard; DNA; 24 BP
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                                                                                                                                                                                                                                                                                                                              99US-00475947.
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                                                                                                                                                                                                                                                                                                                                                                       (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphic probability.
                                                                                                                                                                                                                                                                                                                                                                                                                Wren JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-208818/20.
                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                       31-DEC-1999;
                                                                                                                                                                                                                                                                                                                                31-DEC-1999;
                                                                                                                                                                                                      JS6472154-B1
                                                                                                                                                                                                                                                 29-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                              Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
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Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises introducing into an animal a nucleic acid molecule comprising a germ line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a method of creating an insertional mutation in the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence represents a primer used to characterise the insertion site of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         L1/enhanced green fluorescent protein (EGFP) retrotransposon cassette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 23; DB 1; Length 24;
100.0%; Pred. No. 4.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24 BP; 23 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; SEQ ID NO 151; 102pp; English.
                                                                                                                                                                                                                                                  Deberardinis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAAAAA 2731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP specific SNPE primer SEQ ID 1311.
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ID AAH38515 standard; DNA; 25 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-OCT-2000; 2000WO-US028436.
                                                                                                                                                        28-APR-1997; 97US-00847844.
01-SEP-2000; 2000US-00653812.
                                                                                   09-AUG-2002; 2002US-00216122.
                                                                                                                       95US-0006831P.
96US-00749805.
                                                                                                                                                                                                                                                  Ostertag E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2001 (first entry)
                                                                                                                                                                                                                (UYPE-) UNIV PENNSYLVANIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 into the mouse genome.
                                                                                                                                                                                                                                                                                   WPI; 2003-863454/80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                              specific promoter.
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                                                                                                                                                                                                                                                   Kazazian HH,
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                                                                                                                         16-NOV-1995;
                                                                                                                                         5-NOV-1996;
                                                  26-JUN-2003
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Gaps

Homo sapiens

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(SEQU-) SEQUEL GENETICS INC.
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                                                                                                                                                                                                           Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking conference of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The Oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or identity of a SNP and for genotyping nucleic acid samples, for e.g. to assess by association analysis the genotypic caid samples, for e.g. to assess by association analysis the genotypic trait suspected of being caused by one or more SNPs. Phenotypic traits include disease e.g. agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular dystrophy, familial hypercholesterolaemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic craits also include symptoms of or suscoptibility to multifactorial clisease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arthritis, multiple sclerosis, microorganism. The method is also useful in forensic investigations and microorganism. The method is also useful in forensic investigations and primer specific for a human SNP containing DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide-based cDNA characterisation; haemoglobin alpha 2; human; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 23; DB 1; Length 25; 100.0%; Pred. No. 4.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human haemoglobin alpha 2 transcript (extreme 3' end).
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                                                                                                                                                                                       Claim 1; Page 56; 83pp; English.
                                          (ORCH-) ORCHID BIOSCIENCES INC.
               99US-0160096P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS11744 standard; DNA; 28 BP.
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                                                                     Picoult-Newburg L, Pohl M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                  WPI; 2001-290930/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 23; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200161051-A1.
             15-OCT-1999;
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Matches
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The invention relates to a method of peptide-based transcript or genetic analysis comprising: (a) providing multiple polynucleotides (I) derived from MRNAs from a biological sample, where (I) has homology to a known creference sequence (II); (b) expressing (I); and (C) assessing a physical property of the expressing products to determine the sequences of (I) by comparison with the predicted properties of polypeptides encoded by (II). The method is useful for transcript or genetic analysis, especially conformed is useful for transcript or genetic analysis, especially concleic acid analysis where the method comprises expressing polypeptides or nucleic acid analysis where the method comprises expressing polypeptides concleic acid analysis where characteristic of the nucleic acid molecule. The peptide is considerably smaller than the DNA molecule that encodes it individual amino acids averages about 110 Daltons each whereas the cindividual amino acids averages about 110 Daltons each whereas the individual amino acids averages about 110 Daltons each whereas the composed of combinations of 20 different amino acids instead of combinations of 4 different nucleotides, e.g., two random DNA fragments of identical composition (e.g., with 10 adenines, 15 combinations of 4 different nucleotides, e.g., two random DNA fragments of identical composition (e.g., with 10 adenines, 15 combinations of distinguished on the basis of mass, the present sequence represents the coding sequence of human character the method of the invention are adding sequence of human character the method of the invention and average of the invention and a cannot be the invention and a cannot be asset to the coding sequence of human character the method of the invention and average of the invention and an example to the coding sequence of human character the method of the invention and a cannot of the invention and and a cannot be adding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                               Transcript, genetic, and especially nucleic acid sequence analysis comprises analysis of hybrid peptide products.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 23; DB 1; Length 28; 00.0%; Pred. No. 5.2e+02;
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/*tag= a
/note= "phosphodiester oligonucleotide"

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                                                                                                                                                             Example 11; Page 30; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT93819 standard; DNA; 26 BP.
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Matches 23; Conservative
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WPI; 2001-514778/56.
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modified_base
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24-FEB-1998
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                                                                                                                                                                                                 Novel phosphodiesteric oligonucleotides AAT93811-27 are based on the generic formula, in the 3'-5' or 5'-3' direction: (GaTa')a''-(GbTb')b''-

(GCTC')C''-(GdTd')d''-(GFTe')e''-(GTTE')g''-1', where: N and different from each other; a, b', c', d', e', f', and g' = 0-10, equal or different from each other; a', b', c', d', e', f', and g' = 0-30, equal or different from each other; a', b', c', d', e', f', and g' = 1-

Is different from each other; a', b', c', d', e', f', and g' = 1-

Is dequal or different from each other; The oligonucleotides are believed to selectively bind and sequester some proteins which are essential to the viability and growth of tumours cell ine. They have specific and elective cytotoxic activity against tumour cells, and can be used for treating tumours of the liquid type, in particular of lymphoblastic origin, and of solid type, in particular lymphomas. The present phosphodiester oligonucleotide, at a concentration of 15 micromolar reduced growth of CCRF-CEM tumoural cells by 76%, which is detectable 48 hours after administration. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                New phospho:di:esteric oligo:nucleotide(s) - which exert a specific and selective cytotoxic effect on tumour cells, for treating both solid and
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Compound activity characterisation-related oligonucleotide SeqID1
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phenotypic attribute; candidate medicine; candidate treatment;
multiple biological descriptor; cell marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 22.8; DB 1; Length 26; 22.3%; Pred. No. 5.2e+02; ve 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26 BP; 0 A; 0 C; 2 G; 24 T; 0 U; 0 Other;
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                                                                                                                                                                                 Claim 10; Page 5; 38pp; English.
                                                                                   Quadrifoglio F;
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20-FEB-2001; 2001US-00790214.
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             96WO-EP005388
                                    95IT-MI002539
                                                                                                                                                                                                                                                                                                                                                                                                                                                 92.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 92.3
nes 24; Conservative
                                                                                                           WPI; 1997-319771/29
                                                             SRL.
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                                                                                                                                                          liquid tumours.
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                                                             (SAIC-) SAICOM
                                                                                    Scaggiante B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
             04-DEC-1996;
                                    04-DEC-1995;
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This invention relates to a novel method for the characterisation of the activity of a compound on cell. The method involves receiving images of cells with a cellular activity and images of other cells treated with the compound, quantitatively characterising phenotypic attributes of the cape of cells with a cellular activity to produce a target phenotype for the compound, and comparing the two phenotype for the compound, and comparing the two phenotypes to cher may be useful for characterising cellular activity. The invention cape may be useful for characterising cellular activity. The invention about properties of substances based upon the chiformation about properties of substances based upon the chiformation about properties of substances based upon the cubstances. The invention is also useful for identifying promising candidates in a search for new and better medicines and treatments using multiple biological descriptors from a single cell markers or components.
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                                                          Characterizing cellular activity of compound, by receiving images of cells with known activity and images of cells treated with compound, characterizing phenotypic attributes of images and comparing the
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WPI; 2002-041423/05
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                                                                                                                                                                                             phenotypes.
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AAD34264 standard; DNA; 25 BP.

AAD34264;

the number of mucleotides in the Oilgonnoleotide, and (b) 5' XXNIYZMZ or the number of mucleotide, and as a most as a content of the Oilgonnoleotide, and as a fences the 3' end of the Oilgonnoleotide, and as a content of the Oilgonnoleotide, where X is a Tor modified T nucleotide, NI is a polynucleotide of 5:10 nucleotide, where NI does not include a CG dinucleotide of 5:10 nucleotide, where NI does not include a CG dinucleotide of 5:10 nucleotide, where NI does not include a CG dinucleotide of 5:10 nucleotide, where NI does not include a CG dinucleotide of 5:10 nucleotide, and where NI is a polynucleotide of 5:10 nucleotide, and where NI is a polynucleotide of 5:10 nucleotide in comprising a molecular selected from chemotherapeutic agents, monoclonal antibodies, and anticancer agents; content as a personal content of a polynucleotide of 5:10 nucleotide or a pharmaceutical composition to the with a therapeutic agents, monoclonal antibodies, and anticancer agents; craditorically increased IL-10 expression in a subject composition to the subject in med of increased IL-10 expression relative to IR-alpha expression; (3) a method of inducing an antigen specific regulatory nucleic acid or composition to a subject by administering an immunostimulatory nucleic acid or composition to a subject by administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an aubject by exposing a subject to a self antigen, and administered in an aubject by exposing a subject to a self antigen. and administered in a subject by exposing a subject to a self antigen. and administered in a subject by exposing a subject to a self antigen. and administered in a subject by exposing a subject to a self antigen. and administered in a subject by exposing a subject to a self antigen. and administered in a subject b The invention relates to an oligonucleotide having the formula: (a) 5' XYNIYZNZ 3' where 5' denotes the 5' end of the oligonucleotide and 3' denotes the 3' and of the oligonucleotide, where X is a T or modified T nucleotide, Y is a C or modified C nucleotide, Z is a G or modified G nucleotide, and N1 and N2 are polynucleotides that do not include a CG dinucleotide, where N1 does not include 5' Z nucleotide, and where a 3' polynucleotide consisting of the YZ dinucleotide and the N2 polynucleotide consisting of the YZ dinucleotides that is at most 45% of the number of nucleotides in the oligonucleotide; and (b) 5' XYNIYZNI 3' the number of nucleotides and (b) 5' XYNIYZNI 3' polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autolimune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present 

Sequence 24 BP; 0 A; 1 C; 0 G; 23 T; 0 U; 0 Other;

Gaps .. 0 0.8%; Score 22.4; DB 1; Length 24; 95.8%; Pred. No. 5.3e+02; Indels 1, Pred. No. 5.3e 0; Mismatches Conservative Local Similarity nes 23; Conserv Query Match Best Loca Matches

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The invention relates to sequence determination oligonucleotides for detecting polymorphic sites in a 5' flanking region of cytochrome P450 ab6 (CYP2D6) gene. CYP2D6 enzymes are involved in the metabolism of many different xenobiotics. Human CYP2D6 gene is located on chromosome 22. The oligonucleotides may be used as in situ hybridisation probes, in ligase-based sequenced determination, as components of diagnostic assays, as probes in sequence determination methods based on mismatches, as hybridisation-based diagnostic assays, and as components of diagnostic microarray. CYP2D6 is useful to predict variations in an individual's ability to metabolise certain drugs. The present sequence is a sense oligonucleotide used for detecting of human CYP2D6 gene 5' flanking chromosome 22; ss. Vascular endothelial protein-tyrosine phosphatase; VE-PTP; mouse; human; gene therapy; cytostatic; VE-cadherin; PCR; primer; 88; vascular endothelial-cadherin. New sequence determination oligonucleotides, useful for detecting polymorphic sites in a 5' flanking region of a CYP2D6 gene, as hybridization probes, as components of diagnostic assays, or in ligase-Gaps Human; cytochrome P450 2D6; CYP2D6; enzyme; detection; xenobiotic; ligase-based sequenced determination; drug metabolism; chromosome Human CYP2D6 gene polymorphic site 385 detecting sense 5' oligo. ö Query Match 0.8%; Score 22.4; DB 1; Length 25; Best Local Similarity 95.8%; Pred. No. 5.4e+02; 1; Indels Lewander T, Oliasson E; Sequence 25 BP; 22 A; 2 C; 1 G; 0 T; 0 U; 0 Other; Murine VE-PTP coding sequence PCR primer #2. region single nucleotide polymorphism (SNP) 0; Mismatches 2707 CTAAAAAAAAAAAAAAAAAAAA 2730 1 CCAAAAAAAAAAAAAAAAAAAAA 24 Claim 2; Page 23; 63pp; English. BP. 27-AUG-2001; 2001WO-IB001544. based sequence determination. 30-AUG-2000; 2000GB-00021286. (GEMI-) GEMINI GENOMICS PLC. Andersson MK, AAL57030 standard; DNA; 23 (first entry) (first entry) Conservative WPI; 2002-329785/36. WO200218638-A2. Homo sapiens. Risinger C, 16-JUL-2002 07-MAR-2002. 23; 11-MAR-2004 AAL57030; RESULT 327 AAL57030/c ઠે 유 

DNA

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molecules are inhibitory to cell growth. The molecules AA064709, AA064711 and AAQ64724 all lacked cytotoxicity. In the novel 2-5A-antisense oligonucleotide chimeric molecules, the antisense region targets the chimeric molecule region of RNA to be specifically cleaved and the 2',5'-linked terraadenylate tail activates the 2-5A RNase. Typical applications are treatment of viral infections (esp. for cleaved on the 2' strengenome), cancer; leukaemia, cardiovascular disorders (e.g. restenosis after angioplasty), genetic disorders, osteoarchritis or rheumatoid arthritis. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Specific cleavage of RNA, useful partic. for treating viral infection, cancers, etc. - by using anti-sense oligo:nucleotide coupled to activator of 2-5A dependent RNase.
    = a) and the antisense
/note= "the 2-5A moiety (*tag = a) and the antisense
sequence (*tag = c) are linked by two 1,4-butanediol
molecules linked through phosphodiester bonds"
                                                                                     /*tag= c
/note= "antisense region, complementary to oligo dT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence was used to determine whether 2-5A-antisense chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 5.4e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                             Lesiak K;
                                                                                                                                                                                                                                                                                                                     (USSH ) US DEPT HEALTH & HUMAN SERVICES. (CLEV-) CLEVELAND CLINIC RES INST.
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Best Local Similarity 100."
Marches 22, Conservative
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                                                                               *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-151315/18.
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                                                                 misc feature
                                                                                                                                                                                                                         20-OCT-1993;
                                                                                                                                                                                                                                                              21-OCT-1992;
                                                                                                                                                                                                                                                                                17-SEP-1993;
                                                                                                                                              WO9409129-A2
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                                                                                                                                                                                  28-APR-1994
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AAF17413
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a polypeptide comprising vascular endochelial-protein tyrosine phosphacase (VE-PTP) or its active fragment or effector, or the nucleic acid encoding the polypeptide or its effector, for use in the manufacture of an agent for monitoring or modulating vascular endothelial-protein tyrosine phosphatase (VE-PTP) or its active fragment or effector, or the nucleic acid encoding the polypeptide or its effector, or the nucleic acid encoding the polypeptide or modulating VE-cadherin mediated processes or disorders.

For monitoring or modulating VE-cadherin mediated processes or disorders, e.g., cancer. The present sequence is a PCR primer shown in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                         Use of the polypeptide comprising vascular endothelial-protein tyrosine phosphatase (VE-PTP) or the nucleic acid encoding the polypeptide for the manufacture of an agent for monitoring or modulating VE-cadherin mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1. .4
/*tag= a
/label= 2.,5'-linked_tetraadenylate
/label= "nucleotides linked through phosphodiester bonds
at hydroxyl groups of 2' and 5' carbons"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2',5'-linked tetraadenylate-anti(dT)18 oligonucleotide chimeric mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense; 2',5'-tetraadenylate; 2-5A dependent RNase activator; RNA cleavage; antiviral therapy; chimeric molecule; PKR; protein synthesis regulation; phosphorylation; eIF-2alpha; eukaryotic translation initiation factor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 22.2; DB 1; Length 23; 95.7%; Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                  Golding M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 0 A; 0 C; 0 G; 22 T; 0 U; 1 Other;
                                                                                                                                                                                                                             (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                  Shima DT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВР.
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                                                                                                                                                                                                                                                                  Vestweber D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ64724 standard; cDNA to mRNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 17; Opp; English.
                                                                                                                                                  08-APR-2003; 2003WO-EP003645
                                                                                                                                                                                      08-APR-2002; 2002EP-00007837
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 95.7
Matches 22; Conservative
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                                                                                                                                                                                                                                                                  Deutsch U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
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                                                                     WO2003084565-A2
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misc_feature
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04-JAN-1995
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replication
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ADQ25630/c
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                                                                                          DNAc comprising a promoter P and an L1 cassette sequence having a core retrotransposon element, useful for random insertion of a heterologous or homologous DNA sequence into a cell genome and for correcting genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises introducing into an animal a nucleic acid molecule comprising a germ line specific promoter.
                                                                                                                                                                 The present invention relates to DNA for a promoter and an L1 cassette sequence having a core retrotransposon element. The invention is useful for random insertion of a heterologous or homologous DNA sequence into cell genome, and for correction of a genetic defect in the cell into which the insertion is made. Genetic defects which may be corrected includes cystic fibrosis, mutations in the dystrophin gene, genetic defects associated with blood clotting and other genetic defects
                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mutation generation; transgenic animal; poly A element; non-LTR; retrotransposon; long terminal repeats; L1; EN domain; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; insertional mutation; germ line specific promoter;
                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                        0.8%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 5.4e+02;
tve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       L1 retrotransposon endonuclease cleavage site seg id 94.
                                                                                                                                                                                                                                                    Sequence 22 BP; 22 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                    Boeke JD;
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                                                    Dombroski BA, Kazazian HH,
                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAAA 2730
                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 14; 87pp; English.
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96US-00749805.
97US-00847844.
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 96US-00749805
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15-NOV-1996; 96US-00749805.
28-APR-1997; 97US-00847844.
01-SEP-2000; 2000US-00653812.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-AUG-2002; 2002US-00216122.
                   (UYDE-) UNIV JOHNS HOPKINS (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                         ADF12348 standard; DNA; 22
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                                                                                                                                                                                                                                                                                              22; Conservative
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                                                                      WPI; 2001-060015/07.
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                                                                                                                                                                                                                                                                                   Local Similarity
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 15-NOV-1996;
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                                                  Moran JV,
                                                                                                                                                                                                                                                                                                                                                                                                              ADF12348;
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                                                                                                                                                                                                                                                                         Query Match
                                                                                                                           defects
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Matches
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                                                                                                                      the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence human L1 retrotransposon EN domain.
                                                                                The invention describes a method of creating an insertional mutation in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amplifying RNA sequences, useful in detecting diseases or mutation, comprises synthesizing first strand cDNA, circularizing first strand cDNA, and replicating the circularized cDNA molecules by rolling circle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA sequences. The method involves synthesising first strand cDNA molecules from RNA molecules, circularising the first strand and replicating the circularised first strand cDNA molecules using rolling circle replication. The method is useful for producing nucleic acid molecules corresponding to RNA molecules in an RNA sample, for identifying or analysing and comparing RNA molecules and or sequences expressed in different cells, tissues and or samples. The invention is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to composition and method for amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 5.4e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22 BP; 22 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
Example 2; SEQ ID NO 94; 102pp; English.
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hes 22; Conservative
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(ABAR/) ABARZUA P.
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also useful in detecting disease (e.g. cystic fibrosis, muscular dystrophy or diabetes), mutation detection, gene discovery, gene mapping (molecular haplotyping), agricultural research, and assessment of predisposition for cancers, e.g. prostate, breast, lung, colon or owarian cancer. The present sequence is a junction-specific cDNA primer. This sequence is used to illustrate the method of invention.
                                                                                                                                                                                                                                                                                                                                                                      Human interleukin-6 gene; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
11. .12
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/note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
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                                                                                                                                                                                                                                                                                                                                               Oligomer IL6803 for forming triplex with HUMIL6 target duplex.
                                                                                                               Query Match 0.8%; Score 22; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 5.4e+02; Matches 22; Conservative 0; Mismatches 0; Indels
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note= "o-xyloso dimer synthon linkage"
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                                                                                        Sequence 22 BP; 0 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
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label= inverted_polarity_region
'note= "see comments"
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
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modified_base
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17-APR-1991;
27-SEP-1991;
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08-APR-1991;
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07-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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The synthetic oligomer is capable of forming a triplex at physiological ph with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human interleuwin 6 gene untranslated sequence contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA cumours and inflammation. The triple helices form under mild conditions the major and inflammation. The triple helices form under mild conditions concitions to harsh conditions. The oligomer contains an inverted polarity region formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso (nucleotides have the 3'positions of xylose sugars linked via the o-concitions and interpretation). Two nucleotides are coupled through a xylene residue to consider stable to nuclease activity. The oligomer is able to inhibit conditions as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.) ö Human interleukin-6 gene, herpes simplex, AIDS, modified, HIV, RSV, HPV, malignancy; hepatitis, inflammation, 88. Gaps /mod\_base= OTHER /note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine" ö Oligomer IL6804 for forming triplex with HUMIL6 target duplex. Length 23; 0; Indels 'note= "o-xyloso dimer synthon linkage" base= OTHER == "OTHER= N4 N4 ethanocytosine" Sequence 23 BP; 2 A; 0 C; 0 G; 21 T; 0 U; 0 Other; /\*tag= c /label= inverted\_polarity\_region /note= "see comments" Score 22; DB 1; Le Pred. No. 5.5e+02; 100.0%; Pred. No. 2708 TAAAAAAAAAAAAAAAAA 2729 Location/Qualifiers herpes malignancy and inflammation 23 TAAAAAAAAAAAAAAAA 2 Claim 12; Page 71; 77pp; English. 뭠. 91WO-US008811. 90US-00617907 0.8%; AAQ30431 standard; DNA; 23 b Ω (first entry) /\*tag= a Best Local Similarity 100. Matches 22; Conservative . 12 12. .23 /\*tag= /\*tag= 'note= (revised) DOE, Key modified\_base modified\_base 25-NOV-1991; 23-NOV-1990; misc\_feature misc\_feature WO9209705-A1 11-JUN-1992. 25-MAR-2003 07-DEC-1992 Synthetic. AAQ30431; Query Match AAQ30431 원 8

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The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human interleukin 6 gene untranslated sequence contg. a purine rich sequence contg. a purine rich sequence contg. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of disease characterised by specific DNA duplex targets, e.g. HPV, HER, HIV, hepatitis B, herpes, malignant continuours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. The oligomer contains an inverted polarity region (nucleotides have the algomer contains an inverted polarity region (nucleotides have the algomer contains an avylene residue to contident synthon. This additional modifications may render the oligomer stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                         New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human, MLH1, MSH2; hMLH1; hMSH2; variant gene; diagnosis; HNPCC, hereditary non-polyposis colorectal cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 22; DB 1; Length 23; 100.0%; Pred. No. 5.5e+02; Attive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                          Matteucci MD, Milligan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human MSH2 (hMSH2) intronic sequence SEQ ID NO:126.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 23 BP; 1 A; 1 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2708 TAAAAAAAAAAAAAAAAAA 2729
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                                                                                                                                                                                                                                                                                               Claim 12; Page 71; 77pp; English.
91US-00643382.
91US-00683420.
91US-00686544.
91US-00686547.
91US-00686547.
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Matches 22; Conservative
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                                                                                                                                                       Froehler B, Krawczyk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ROBB/) ROBBINS D.
(LING/) LIN-GOERKE J L.
                                                                                                                    (GILE-) GILEAD SCI INC
                                                                                                                                                                                           WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JS2001044936-A1.
                08-APR-1991;
17-APR-1991;
17-APR-1991;
                                                                  17-APR-1991;
27-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-1999;
   18-JAN-1991;
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The present invention describes a variant human MLH1 or MSH2 gene. Also described are: (1) a method for diagnosing or predicting susceptibility to hereditary non-polyposis colorectal cancer (HNPCC), comprising cereahidates presence of, or susceptibility to HNPCC; (2) a method of identifying mutants in splice donor or acceptor sites of a human MLH1 gene, comprising sequencing splice donor or acceptor sites of the gene with intronic primers for the human MLH1 gene and analysing the sequence to identify any mutants; (3) a method of identifying mutants in splice donor or acceptor sites of a human MSH2 gene, comprising sequencing splice donor or acceptor sites of the gene with intronic primers for the human MSH2 gene and analysing the sequence to identify any mutants; (3) a method of identifying mutants in splice donor or acceptor sites of the gene with intronic primers for the human MSH2 gene and analysing the sequence to identify any mutants; and (4) a transgenic model system for colorectal cancer comprising cells expressing the variant MLH1 or MSH2 gene. The hMLH1 and hMSH2 variants are used to diagnose or determine a patients

CC susceptibility to hereditary non-polyposis colorectal cancer. ABL01648 to ABL01745 and ABL01746 to ABL01831 represent hMLH1 and hMSH2 gene fragments from the present invention and the present invention and the present invention and the present invention of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           production; venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RT-PCR primer used for cDNA synthesis from scorpion toxin RNA Seg 285.
                                                                                                              New variants of the human MLH1 and MSH2 genes for diagnosing or determining a predisposition for hereditary non-polyposis colorectal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 22; DB 1; Length 23; 100.0%; Pred. No. 5.5e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RT-PCR; 88; toxin; immunogenicity; antigen; antibody p vaccine; diagnosis; primer; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23 BP; 21 A; 0 C; 1 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYME-) UNIV MEXICO NACIONAL AUTONOMA.
(SILA-) LAB SILANES SA DE CV.
                                      Ling JC;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-NOV-2003; 2003US-00721793
                                      D, Lin-Goerke JL,
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                                                                           WPI; 2002-105577/14.
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(LING/) LING J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2005065331-A1.
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Possani PLD;
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                                    Robbins
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Novel isolated scorpion toxin polypeptide, useful for preventing envenomation from scorpion stings, and as vaccine to prevent envenomation from venom of scorpions of genus Centruroides.

Disclosure, SEQ ID NO 285; 135pp; English

This invention relates to novel scorpion toxin polynucleotides and the encoded proteins thereof having any one of 142 fully defined amino acid sequences given in the specification. Specifically, it refers to long chain toxins that block the sodium channels of excitable cells and also chain toxins that affect Erg type potassium channels. The present invention describes immunogenic or antigenic compositions comprising a scorpion toxin protein or fragment thereof, which can be used for the can revent entibodies that are able to bind to and neutralize the in scorpion toxin protein or fragment thereof, which can be used for the compositions or appropriate recombinant fusion proteins in the development of vaccines that can prevent envenomation from stings of scorpions of the genus centruroides. Furthermore, it provides a diagnostic method that uses an immunogenic matrix for the purification of specific immunoglobulins present in a sample that can determine the species of scorpion that has tunny an individual through the detection of identifying antibodies. In addition, it provides methods that are useful for treating envenomation from scorpion stings. This oligonucleotide is an RT-PCR primer used to synthesize cDNA from scorpion toxin total RNA of the invention.

Sequence 24 BP; 0 A; 0 C; 0 G; 22 T; 0 U; 2 Other;

Gaps ö Score 22; DB 1; Length 24; Pred. No. 5.6e+02; 0; Indels 100.0%; Preu. ....ive 0; Mismatches 0.8%; Conservative Query Match Best Local Similarity Matches 22; Conserv

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ADF12409 standard; DNA; 26 RESULT 336 ADF12409 

ADF12409;

(first entry) 12-FEB-2004 L1 retrotransposon endonuclease cleavage site #1.

gene therapy; insertional mutation; germ line specific promoter; mutation generation; transgenic animal; poly A element; non-LTR; retrotransposon; long terminal repeats; L1; EN domain; endonuclease; cleavage site; ds

Homo sapiens

US2003121063-A1.

26-JUN-2003

09-AUG-2002; 2002US-00216122

95US-0006831P. 96US-00749805. 97US-00847844. 16-NOV-1995; 15-NOV-1996; 28-APR-1997

01-SEP-2000; 2000US-00653812

Deberardinis Ostertag E, Kazazian HH,

(UYPE-) UNIV PENNSYLVANIA.

WPI; 2003-863454/80.

Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises

introducing into an animal a nucleic acid molecule comprising a germ line specific promoter.

Example 2; Fig 14A; 102pp; English.

The invention describes a method of creating an insertional mutation in the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence trapresents an examplary cleavage site of the endonuclease encoded by human L1 retrotransposon EN domain. 

Sequence 26 BP; 22 A; 0 C; 0 G; 0 T; 0 U; 4 Other;

ö Gaps ö 0.8%; Score 22; DB 1; Length 26; 100.0%; Pred. No. 5.88+02; tive 0; Mismatches 0; Indels 22; Conservative Query Match Best Local Similarity Matches

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RESULT 337

BP. ABK86170 standard; DNA; 25 ABK86170

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ABK86170;

(first entry) 24-SEP-2002 Oligo dT primer #3 used in method to study gene expression.

Oligo dT primer; gene expression analysis; primer; ss.

Synthetic.

WO200236828-A2.

10-MAY-2002.

01-NOV-2001; 2001WO-US045401.

01-NOV-2000; 2000US-0244933P.

(GENO-) GENOMIC SOLUTIONS INC.

Nagel AC; Kane MD, Dombkowski AA,

WPI; 2002-508123/54.

Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identimer having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.

Example 2; Page 21; 45pp; English.

The invention relates to systems for identification and characterisation of gene expression in one or more samples, comprising an identimer having a specific oligo-dT primer sequence, where the identimer comprises a detectable marker at its 5' end. The system is useful for identifying any or all genes expressed in a given in vivo or in vitro RNA sample, as well as the relative differences in mRNA between 2 or more samples, where that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising compositions and systems that incorporate new strategies where molecular

or biochemical assay compositions and systems are linked to DNA or RNA esquence databases for optimal resource efficiency in assaying gene expression. The system has the following advantages over existing methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence information in addition to information concerning changes or differences in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates CDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology techniques; and (d) does not require prior knowledge of the sequence of the genome of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present sequence represents an oligo dT primer used in the method of the invention

Sequence 25 BP; 0 A; 0 C; 2 G; 23 T; 0 U; 0 Other;

Score 21.8; DB 1; Length 25; Pred. No. 5.9e+02; 0; Mismatches 2; Indels 2709 AAAAAAAAAAAAAAAAAAAA 2733 25 AAACCAAAAAAAAAAAAAAA 1 0.8%; 23; Conservative Best Local Similarity Matches 23; Conserv Query Match 셤 ઠે

ADO81056 standard; DNA; 25 BP 29-JUL-2004 (first entry) ADO81056; RESULT 338 ADO81056/ 

gene typing; polymorphic microsatellite loci; PML; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow; microsatellite; PCR; primer; ss. Cow prion protein microsatellite locus primer #68.

Bos taurus.

DE10236711-A1

26-FEB-2004

09-AUG-2002; 2002DE-01036711.

09-AUG-2002; 2002DE-01036711

(UYHO-) UNIV HOHENHEIM

Geldermann H, Preuss S,

WPI; 2004-215730/21.

Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.

Example 3; Page 28; 64pp; German.

The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML, and predisposis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a

disease-related gene, that are associated with a predisposition to diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.

888888888

Sequence 25 BP; 0 A; 2 C; 0 G; 23 T; 0 U; 0 Other;

Gaps ö DB 1; Length 25; Indels 2, 0.8%; Score 21.8; DB 1; 92.0%; Pred. No. 5.9e+02; iive 0; Mismatches 2 2709 AAAAAAAAAAAAAAAAAAAAAAA 2733 23; Conservative Similarity Query Match Best Local { Matches 8

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25 AAAAGGAAAAAAAAAAAAAAAA 1

BP. ADO81061 standard; DNA; 25 (first entry) 29-JUL-2004 AD081061; AD081061 

RESULT 339

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Gaps

.; 0

Cow prion protein microsatellite locus primer #73.

gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow; microsatellite; PCR; primer; ss.

Bos taurus.

DE10236711-A1.

26-FEB-2004.

09-AUG-2002; 2002DE-01036711.

09-AUG-2002; 2002DE-01036711.

(UYHO-) UNIV HOHENHEIM.

Geldermann H, Preuss

Han Y;

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WPI; 2004-215730/21.

Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons

Example 3; Page 28; 64pp; German.

as template a DNA sample containing at least one segment of (1); and determining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and predisposis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a disease related gene, that are associated with a predisposition to diseases and for predisgnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormmones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (I) that includes PML, using

ngs.res

AAF16627;

Gastric acid production inhibiting oligonucleotide SEQ ID NO: 114.

(first entry)

13-MAR-2001

Gastric acid disturbance; gastric reflux; gastritis; dyspepsia; stomach ulcer; duodenal ulcer; Helicobacter pylori; antisense; DNA-RNA hybrid; ss.

Homo sapiens.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides oligonucleotides, and methods for their use, which are useful in modulating the action of proteins involved in gastric acid production. The target protein is preferably the histamine H2 receptor or one of the proteins which form part of the gastric proton pump. The sequences and methods of the invention are useful in the treatment of gastric reflux, gastritis, dyspopsia, stomach ulcers, duodenal ulcers and other gastric acid disturbances, most of which are caused by Helicobacter pylori
sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating gastric acid disturbance by administering an oligonucleotide which modulates the activity of a polypeptide involved in gastric acid production or secretion.
                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                          Gastric acid production inhibiting oligonucleotide SEQ ID NO: 103
                                                                                                                                                                                                                                                                                                                                                                      Gastric acid disturbance, gastric reflux, gastritis, dyspepsia, stomach ulcer, duodenal ulcer, Helicobacter pylori, antisense; DNA-RNA hybrid; ss.
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6e+02;
                                                                          Score 21.8; DB 1; Length 25;
Pred. No. 5.9e+02;
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                                              Sequence 25 BP; 0 A; 2 C; 0 G; 23 T; 0 U; 0 Other;
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                                                                                                          0; Mismatches
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                                                                            0.8%;
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                                                                                                                                                                                                                                                  AAF16616 standard; DNA; 26
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Best Local Similarity 92.07
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The present invention provides oligonucleotides, and methods for their use, which are useful in modulating the action of proteins involved in gastric acid production. The target protein is preferably the histamine H2 receptor or one of the proteins which form part of the gastric proton pump. The sequences and methods of the invention are useful in the tracement of gastric reflux, gastritis, dyspepsia, stomach ulcers, duodenal ulcers and other gastric acid disturbances, most of which are caused by Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                  Treating gastric acid disturbance by administering an oligonucleotide which modulates the activity of a polypeptide involved in gastric acid production or secretion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 21.4; DB 1; Length 23; 95.7%; Pred. No. 6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      electrophoresis apparatus; variant gene isolation; gene mutation detection; variant gene detection; single nucleotide polymorphism analysis; SNP detection; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 1 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                       Example 3; Page 152; 164pp; English.
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                                                                                                                                                                                                                                    24-MAY-1999;
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AAF16627 standard; DNA; 23

RESULT 341 AAF16627/ ID AAF1

invention comprises an electrophoresis apparatus for isolating

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention comprises an electrophoresis apparatus for isolating variant genes. The apiarists consists of: a sealed flow path filled with buffer containing linear polymer and DNA joint controlling agent, and containing DNA conjugate for separation, DNA conjugate for purification, and DNA conjugate for assay; and a heating apparatus for heating the portion of sealed flow path in which the DNA conjugate for assay is fixed. The electrophoresis apparatus is useful for isolating a variant gene. The electrophoresis apparatus is also useful in gene diagnosis for detecting the presence or absence of a gene mutation, or variant gene, and for single nucleotide polymorphism analysis. The present DNA sequence was shown in a figure exemplifying the method of the invention.
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                                                                                                                                                Electrophoresis apparatus useful for isolating variant gene, comprises heating apparatus, and sealed flow path comprising linear polymer and DNA joint controlling agent, with DNA conjugates for separation, purification and assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Electrophoresis apparatus useful for isolating variant gene, comprises heating apparatus, and sealed flow path comprising linear polymer and DNA joint controlling agent, with DNA conjugates for separation, purification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene mutation detection; variant gene detection;
single nucleotide polymorphism analysis; SNP detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 23 BP; 22 A; 0 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          electrophoresis apparatus; variant gene isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Electrophoresis apparatus-related DNA sequence #3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAAAAAAAAAAA 2731
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                                                                       (MATU ) MATSUSHITA DENKI SANGYO KK
                                                                                                                                                                                                                                               Disclosure; Fig 2; 21pp; Japanese.
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28-MAR-2003; 2003JP-00091194
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                                     28-MAR-2003; 2003JP-00091194
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ADT55095 standard; DNA; 23
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Best Local Similarity 95.7
Matches 22; Conservative
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and as
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Disclosure; Fig 4; 21pp; Japanese

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variant genes. The apiarists consists of: a sealed flow path filled with buffer containing linear polymer and DNA joint controlling agent, and containing DNA conjugate for separation, DNA conjugate for purification, and DNA conjugate for assay; and eating apparatus for heating the portion of sealed flow path in which the DNA conjugate for assay is fixed. The electrophoresis apparatus is also useful for isolating a variant gene. The electrophoresis apparatus is also useful in gene diagnosis for detecting the presence or absence of a gene mutation, or variant gene, and for single nucleotide polymorphism analysis. The present DNA sequence was shown in a figure exemplifying the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 21.4; DB 1; Length 23; 95.7%; Pred. No. 6e+02; Live 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human phosphatidylinositol-3 kinase 35 cDNA PCR primer #2.
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                                                                                                                                                                                                                                                                                                                                               Sequence 23 BP; 0 A; 1 C; 0 G; 22 T; 0 U; 0 Other;
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hes 22; Conservative
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AAI66361/c
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Matches
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Crompton AM;

Vaisberg EA,

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This invention relates to a novel method for the characterisation of the activity of a compound on cell. The method involves receiving images of cells with a cellular activity and images of other cells treated with the compound, quantitatively characterising phenotypic attributes of the image of cells with a cellular activity to produce a target phenotype for the cellular activity to produce a target phenotype for the compound, and comparing the two phenotype for determine whether the compound possesses cellular activity. The invention can be useful for characterising cellular activity. The invention about properties of substances based upon the information about properties of substances based upon the information is also useful for identifying promising candidates in a search for new and better medicines and treatments using multiple biological descriptors from a single cell markers or components.
                                                                                                                                                                Characterizing cellular activity of compound, by receiving images of cells with known activity and images of cells treated with compound, characterizing phenotypic attributes of images and comparing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Compound activity characterisation-related oligonucleotide SeqID2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate treatment; multiple biological descriptor; cell marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24 BP; 0 A; 1 C; 1 G; 22 T; 0 U; 0 Other;
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Pred. No. 6.1e+02;
0; Mismatches 1;
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                                                                                            Adams CL,
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20-FEB-2001; 2001US-00790214.
26-APR-2000; 2000US-0199778P.
20-FEB-2001; 2001US-00790214.
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                                                                                              Sabry JH,
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                                                         (CYTO-) CYTOKINETICS INC
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                                                                                                                                   WPI; 2002-041423/05.
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Best Local Similarity
                                                                                              Oestreicher DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention provides the protein and coding sequences of human cyclophilin-40-12.54. The sequences can be used in the treatment of immunopathy and cancer. The present sequence is a PCR primer for the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   olypeptide-human cyclophilin-40-12.54 and polynucleotide for coding it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                 cyclophilin-40-12.54; immunopathy; cancer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Compound activity characterisation-related oligonucleotide SegID6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate treatment; multiple biological descriptor; cell marker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 21.4; DB 1; Length 24; 95.7%; Pred. No. 6.1e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                            Human cyclophilin-40-12-54 coding sequence PCR primer #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 17 (Disclosure); 33pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BODE-) BODE GENE DEV CO LTD SHANGHAI.
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                                                                                                                                                                                                                                                                                                                                                                                                 28-JUN-2000; 2000CN-00116823
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ADG16131 standard; DNA; 24
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                                                                                                                                                                         (first entry)
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                                                                                                  AAL47515 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-305482/35.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                       Ното варіеня
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                                                                                                                                                                         13-SEP-2002
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         23
                                                                                                                                      AAL47515;
                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 346
ADG16131/c
                                                               RESULT 345
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Length 24; Indels Crompton AM;

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This invention relates to a novel method for the characterisation of the activity of a compound on cell. The method involves receiving images of cells with a cellular activity and images of other cells treated with the compound, quantitatively characterising phenotypic attributes of the image of cells with a cellular activity to produce a target phenotype for the collular activity to produce a target phenotype for the compound, and comparing the two phenotypes to the cellular activity of a compound, for determine whether the compound possesses cellular activity. The invention may be useful for characterising cellular activity of a compound, for cherming information about properties of substances based upon the information about properties of substances based upon the information about properties of substances based upon the conformation about of living or non-living cells exposed to substances. The invention is also useful for identifying promising candidates in a search for new and better medicines and treatments using multiple biological descriptors from a single cell markers or components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   alysis of cDNA and gene expression - by amplification of mRNA followed digestion with restriction enzymes.
  Characterizing cellular activity of compound, by receiving images of cells with known activity and images of cells treated with compound, characterizing phenotypic attributes of images and comparing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 21.4; DB 1; Length 24; 95.7%; Pred. No. 6.1e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 2 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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                                                                                                         Disclosure; Fig 18; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ75712 standard; DNA; 21 BP.
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hes 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis of
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mammaglobin antigen, then returned to the patient. Expression of mammaglobin is elevated in 27% of stage I primary breast cancers, so it represents a marker useful for diagnosis of this disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present primer was used to reverse transcribe RNA encoding a human
                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                          Human; mammary-specific protein; mammaglobin; antigen; vaccine;
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                                                                                                                                                                                                                                                                                                                                        mammaglobin-expressing cancer; breast cancer; autologous tumor lymphocyte; diagnosis; marker; primer; ss.
                             Score 21; DB 1; Length 21;
Pred. No. 6.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; tive 0; Mismatches 0; Indels
                                                           0; Indels
Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                           Primer used to reverse transcribe mammaglobin RNA
                 0.8%; Scor.
100.0%; Pred. No. c.
                                                                                        2705 TACTAAAAAAAAAAAAA 2725
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                                                                                                                     21 TACTAAAAAAAAAAAAA 1
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                                                                                                                                                                                                 AAX26973 standard; cDNA; 21 BP.
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                                                                                                                                                                                                                                                             (first entry)
                                                          21; Conservative
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                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-SEP-1998;
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                                                                                                                                                                                                                                                             25-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                AAX26973;
                             Query Match
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                                             Local
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AAZ44350/c
ID AAZ4431
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                                                           Matches
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AAX26973/
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(COLE-) COLEY PHARM GMBH.
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                                                                                                                                                                                                                                                                                                    This invention describes novel purified aptameric oligonucleotides which have antimicrobial, cytostatic and immunosuppressive activity. The oligonucleotides are useful for binding to and preventing or inhibiting the biological function of a protein kinase or a target molecule and for detecting the presence or absence of a target molecule in biological samples. The oligonucleotides are also useful for prophylactic and therapeutic treatment of diseases such as cancer, autoimmune diseases and diseases caused by pathogenic microorganisms. This sequence represents a primer used in the method of the invention
                                                                                                                                                                                                                                                     Oligonucleotides inhibiting protein kinase, useful for treating diseases such as cancer and autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            natural antisense mRNA enrichment; antisense-based therapy; primer; ku autoantigen; ss.
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                                                        Antimicrobial; cytostatic; immunosuppressive; protein kinase; prophylactic; therapy; treatment; cancer; autoimmune disease; pathogenic microorganism; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.8%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 6.1e+02;
Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human ku autoantigen amplifying KU_FOR primer.
                                                                                                                                                                                                 (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                        Protein kinase inhibiting primer #12.
                                                                                                                                                                                                                                                                                      Example 8; Col 27-28; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAAAA 2729
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                                                                                                                                                           95US-00416214
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD03631 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                      (first entry)
                                                                                                                                                                                                                    Bergan R, Neckers L;
                                                                                                                                                                                                                                      WPI; 2000-104623/09.
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                                                                                                                                                           04-APR-1995;
                                                                                                                                                                              04-APR-1995;
                                                                                                 Unidentified
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                     04-APR-2000
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                                                                                                                     JS5998596-A
                                                                                                                                         07-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAD03631;
    AAZ44350;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human;
RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 351
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The invention relates to a method for enrichment of natural antisense messenger RNA. This method involves generating a population of cDNA from messenger RNA. This method involves generating a population of cDNA from melecules using the generated cDNA to produce double stranded hybrid concludes using DNA polymerase with an exonuclease activity, amplifying concludes using DNA polymerase with an exonuclease activity, amplifying the double stranded molecule using a deoxythymidine (dT) primer and connict the amplified double stranded molecule. This method is useful for enrichment of natural antisense mRNA from any natural source of RNA. It is used to detect whether mRNAs have a natural anti-sense counterpart. The method provides a basis for finding new genes with important cellular regulatory roles or new regulatory information for known genes and provides a starting material for development of an antisense-based provides a starting material for development of an antisense-based the same gene or transcript is involved. The present sequence is KU FOR reverse transcription PCR (RT-PCR) primer used in midogenous antisense mann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccine, cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         autoantigen sequence. This primer is used in endogenous antisense identification (EASI) procedure for enrichment of natural antisense mRNA
                                                                                                                                                                                                                                                                                          Enrichment and detection of natural antisense mRNA comprises generating double stranded hybrid cDNA using a polymerase with an exonuclease activity, amplifying using a DT primer and cloning.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 6 A; 4 C; 4 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example, Page 12; 37pp; English.
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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99US-0157843P
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                                                                            (QUAR-) QUARK BIOTECH INC
                                                                                                                                                                                                                             WPI; 2001-266326/27.
                                                                                                                                                     Einat P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200122972-A2
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    06-OCT-1999;
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Schmidt W, Hiller R,
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                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxovitidae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), tungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                            ö
                                          Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                   0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide used to produce branched chain compounds.
                                                                                                                                                                                                                                          present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                               Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
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"NH2-C6 attached"
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note= "NH2-C6 attached"
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"branch present"
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                                                                         Claim 101; Page 56; 338pp; English.
 Vollmer J;
                                                                                                                                                                                                                                                                                                                                                   21 AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                       AAH42480 standard; DNA; 21 BP.
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                                                                                                                                                                                                                                                                                                         21; Conservative
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/*tag=
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 Schetter C,
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AAH42480/c
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molecies optionally extended by a polymerase. The branched chain compounds of the invention are used in nucleic acid synthesis reaction, primer extension reaction, reverse transcription reaction of RNA into DNA, nucleic acid hybridization experiment (for identifying sequence of nucleic acid), and nucleic acid amplification experiment (for analysing the expression pattern of genes). The compounds are also used in solidible invention to produce branched chain compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method of amplifying nucleic acids, involving forming a single-stranded DNA to an mRNA in a sample with a primer, synthesising a DNA strand complementary to the single-stranded DNA to form a double-stranded DNA, adding a single or double-stranded adapter DNA to the double-stranded DNA, adding a single or double-stranded adapter DNA to the double-stranded DNA, and amplifying the DNA strand using a second primer with a nucleic acid sequence in the adapter DNA. This can be used to amplify nucleic acids to monitor mRNA expression, which is applicable in producing e.g. CRNA libraries, CDNA libraries, DNA
                                                                                                                                                                                                                                                                                                                     The specification describes branched compounds containing nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amplifying nucleic acids with base sequences of mRNAs in sample while sustaining the ratio among them used to monitor mRNA expression, applicable in producing e.g. CRNA library and DNA microarrays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                      Branched compounds useful in e.g. nucleic acid synthesis reaction comprises nucleic acid moieties optionally extended by a polymerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid amplification; adapter DNA; human; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 6.1e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human KUAPP70 gene PCR primer SEQ ID NO: 40.
   Mueller M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAA 2729
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                                                                                                                                                                                                                                                      Example 1; Page 10; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Huber M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH45788 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                      WPI; 2001-466959/51.
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Page 172

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The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; procancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; Blaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
microarrays or membrane arrays in gene engineering and gene expression analysis, and in drug development and health maintenance and management. The present sequence is a PCR primer described in the exemplification of the invention
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                                                                                                                                    DB 1; Length 21;
6.1e+02;
hes 0; Indels
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                                                                                                                                    0.8%; Score 21; DB 100.0%; Pred. No. 6.1
                                                                                                                                                                                                              2416 TTACGGGCTGAAGAGTGGTCT 2436
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                                                                                                                                                                                                                                        TTACGGGCTGAAGAGTGGTCT
                                                                                                                                                                                                                                                                                                                                                ₽.
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                                                                                                  Sequence 21 BP; 6 A; 8 C; 3
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                                                                                                                                                     Local Similarity 100.
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The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigons and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, concert, concert, concert, concert, connective tissue cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, insequencer, luyer cancer, luyer cancer, luyer cancer, lug cancer, lug cancer, lumphoma, non-CC cancer, pancreatic cancer, myeloma, oral cavity cancer, ovarian CC cancer, pancreatic cancer, prostate cancer, tabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                     Gaps
                                                                                                                                                                                                                                                                                                             Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbone; ss.
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Score 21; DB 1; Length 21;
Pred. No. 6.1e+02;
0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                  Immunostimulatory nucleic acid SEQ ID NO: 840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 309; 312pp; English.
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        0.8%;
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                                      21; Conservative
                                                                                                                                                                                    ABL39404 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hartmann G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-154611/20.
        Query Match
Best Local Similarity
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100.0%; Pred. ...

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Matches

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Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;

Local Similarity

Query Match

2709 AAAAAAAAAAAAAAAAAA 2729

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The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contect dermatisis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                         Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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                                                                                                                              Immunostimulatory nucleic acid #881.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2001; 2001US-0279642P
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                                                                               25-SEP-2003 (first entry)
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1es 21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Krieg AM, Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KRIE/) KRIEG A M.
(BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                             US2003050268-A1.
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                                                                                                                                                                                                                                                                                             Synthetic.
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                                  ACH03246;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             performance animal. The method involves determining in sample abundance of expressed target nucleic acid, transmitting digital sample signal to remote diagnostic server; processing digital sample signal at remotely located diagnostic server; processing digital sample signal to remote diagnostic server; processing digital sample signal to remotely located diabase to correlate digital signal with digital information and returning report of particular condition of animal. The method is useful for assessing a condition of a performance animal preferably human, dog or camel. The condition can be an athletic ability and a condition that enhances, hinders, impedes or does not change an expected ability of the performance animal; and also normal, pre-clinical, overt progress and/or stages of disease, undiagnosed of unclassified conditions, presence of drugs, response to environmental conditions. Diseases assessed by the invention include laminitis, lameless, viral or bacterial disease, of gastric ulders, respiratory aliments, fractures, epistaxis, musculoskeletal damage or disorders and joint diseases. The present sequence is a primer used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states, of performance animal involves analyzing nucleic acid expression.
                                                                                                                                                                                                                                                                                                                                   Laminitis, viral disease, vaccine, bacterial disease, primer, epistaxis, gastritis, gastric ulcer, respiratory ailment, fracture, joint disease, musculoskeletal damage, ss.
                                                                                                                                                                                                                                                                                      Regular oligo dT primer used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENO-) GENOMICS RES PARTNERS PTY LID.
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                           21 AAAAAAAAAAAAAAAAAAAA
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03-MAY-2002; 2002WO-AU000553.
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29-JUN-2001; 2001US-00896941
                                                                                                                                                AAD51323 standard; DNA; 21
                                                                                                                                                                                                                                            (first entry)
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Best Local Similarity 100.(
Matches 21, Conservative
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WO200290579-A1.

14-NOV-2002.

Brandon RB;

Unidentified

16-APR-2003

AAD51323;

RESULT 357

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AAD51323

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Gaps

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0; Indels

08-MAY-2003

ACH03246 standard; DNA; 21 BP

RESULT 358 ACH03246/c ID ACH03;

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RESULT 361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method for the amplification of DNA that maintains genes and copy number of the sequence. This method is useful for easy and operable amplification of DNA. The method was carried out by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amplification of DNA maintaining genes and copy number of the sequence on a genome, and their ratios in the resultant DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer for amplifying the ATP dependant DNA helicase gene #SEQ ID 69.
                                                                                                                                                                                                                                                                            The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnoshing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                     g and/or preventing allergy or asthma using an immunostimulatory acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA amplification; copy number; polymerase chain reaction; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 21; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 6.1e+02; Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KOKU-) KOKURITSU GAN CENT SOCHO.
(IYAK-) IYAKUHIN FUKUSAYO HIGAI KYUSAI KENKYU SH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; SEQ ID NO 69; 33pp; Japanese.
                                                                                                                                           Fouron Y;
                                                                                                                                                                                                                                                     Disclosure; Page 17; 221pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAY-2001; 2001JP-00137858.
                02-FEB-2001; 2001US-00776479
                                              03-FEB-2000; 2000US-0179991P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC24379 standard; DNA; 21
                                                                                                                                           Bratzler RL, Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                         WPI; 2003-657977/62.
                                                                             BRATZLER R
PETERSEN D
                                                                            (BRAT/) BRATZLER F
(PETE/) PETERSEN I
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP2002345466-A.
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                                                                                                                                                                                                                       nucleic
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nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a S'-invariable anchor region, the complement of which is present at chart comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) modecular filaments and binding of single
                                                                                                                                                                                                                                                                                                                           ö
                                       The current sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                priminally standard to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
fragmentation genomic DNA, preparation of blunt end of the fragmented DNA, ligation of an adapter to the blunted DNA, PCR of the ligated DNA 2 steps, and confirmation of the amplified APC gene. The current sequen represents a PCR primer used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                           0.8%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                     Sequence 21 BP; 6 A; 8 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat DNA microarray capture oligonucleotide #64.
                                                                                                                                                                                                                                                                                                                                                                                                   2436
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADK01344 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                           21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hybridisation;
od; nerve; germ
                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   blood; nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004
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Best Local S
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acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;

Score 21; DB 1; Length 21; Pred. No. 6.1e+02; 0; Indels 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAAA 2729 21 AAAAAAAAAAAAAAA 1 Matches 21; Conservative Best Local Similarity Query Match ò 셤

ADK01341 standard; DNA; 21 BP. 06-MAY-2004 (first entry) ADK01341; ADK01341 

Rat DNA microarray capture oligonucleotide #61.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Dieck HT, Hoppe H; Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable constant regions.

Example; Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using applure agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable and non-variable regions. The capture oligonucleotides at cast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-90, particularly 15-25, T residues. The capture capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular sieve.

and hair root, bload, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cells, and etect very small differences in the nucleic acid pool. Since the method as based on comparison of nucleic acid pool. Since the genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. especially in mucosal, The method is used (i) for analysis of patterns,

Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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Gaps ö

Gaps ö Length 21; 0; Indels Score 21; DB 1; Le Pred. No. 6.1e+02; 0.8%; Scc. 100.0%; Pred. No. c. 0; Mismatches Conservative Similarity 21; Query Match Best Local S Matches

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ВР ADK01330 standard; DNA; 21 RESULT 363 ADK01330/

ADK01330; 

(first entry) 06-MAY-2004

Rat DNA microarray capture oligonucleotide #50.

mucosal; hair root; ss; hybridisation; capture oligonucleotide; pattern; muc blood; nerve; germ cell; food additive; food supplement

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Boekenkamp D, Dieck HT,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded mucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5' invariable and non-variable regions. The capture oligonucleotides have a 5' invariable and non-variable complement of which is present at least once in each nucleic acid and a 3' variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorte of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-

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conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and inxrures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acid from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADX01281-ADX01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
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0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; tive 0; Mismatches 0; Indels 2707 CTAAAAAAAAAAAAAAA 2727 21 CTAAAAAAAAAAAAAAAA 1 Local Similarity 100. ses 21; Conservative Query Match Best Loca Matches

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Rat DNA microarray capture oligonucleotide #8. ADK01288 standard; DNA; 21 BP (first entry) 06-MAY-2004 ADK01288; RESULT 364

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. 28-FEB-2002; 2002DE-01008794. DE10208794-A1. 04-SEP-2003. Rattus sp. 

Boekenkamp D, Dieck HT, Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH.

28-FEB-2002; 2002DE-01008794.

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region

comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic additives or supplements, especially minerals, trace elements, organic mixtures. The method provides rapid, inexpensive and reproducible cailows inaging of the complete pattern of all nucleic acid side from cells. It can detect very small differences in the nucleic acid side in a cell, and can anticative in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual control of the invention is possible. ABK01281-ABK01344 represent capture probes used in the method of the invention. 

Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;

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Gaps .; 0

Gaps ; 0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; tive 0; Mismatches 0; Indels 21; Conservative Query Match Best Local Similarity Matches 21; Conserv

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Human ATP5F1 gene, RT-PCR primer #1. BP. ADM96310 standard; DNA; 21 17-JUN-2004 (first entry) ADM96310; RESULT 365 ADM96310/

ss; human; H+ transporting; mitochondrial ATP synthase; subunit B; isoform 1; ATP5F1; reverse transcriptase; RT-PCR; primer. US2003211483-A1. Synthetic

13-NOV-2003.

Schroth GP; 09-MAY-2002; 2002US-00144179. 09-MAY-2002; 2002US-00144179. Schroeder BG, Chen C, (SCHR/) SCHROEDER B G. (SCHR/) SCHROTH G P. CHEN C. (CHEN/) 

Enriching low abundance polynuclectides in a sample, useful for gene expression analysis, comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer to block polymerase activity on high WPI; 2003-901581/82.

Example 1; Page 20; 43pp; English.

abundance species.

The invention relates to a method of enriching a low abundance polynuclectide in a sample of polynuclectides comprising a low abundance and a high abundance polynuclectide. The method comprises exposing the

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present sequence represents a reverse transcriptase (RT)-PCR primer which was used to amplify the human import precursor of subunit B of the H+ transporting, mitochondrial ATP synthase, subunit B, isoform 1 (ATPSF1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Selective enrichment; gene expression; RT; reverse transcriptase; primer;
                                                                                                                                                                                                                                                         methods are useful in selective enrichment of low abundance polynucleotides in a sample. The pool of enriched polynucleotides may be used in analysing gene expression and in creating CDNA libraries. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RT primer used in the synthesis of an artificial gene transcript.
                                                                                                                                                                                                                                                                                                                                         0.8%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 6.1e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ88057 standard; DNA; 21 BP.
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                                                                                                                                                                                                                                                                                                                                           Query Match
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ADJ88057/c
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The present invention relates to methods for the selective enrichment of low abundance polynucleotides. The invention is useful for analysing gene expression in a sample and creating cDNA libraries. The present sequence is reverse transcriptase (RT) primer used in the synthesis of an
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                                                                                                         Enriching low abundance polynucleotide relative to a high abundance polynucleotide in a sample, for analyzing gene expression and creating cDNA libraries, comprises blocking polymerase activity on high abundance
                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Double-stranded cDNA synthesis; cDNA first strand synthesis; cDNA second strand synthesis; RNA template; RNA amplification; differential gene expression; primer; ss.
                                                                                                                                                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Control primer used in cDNA first strand synthesis.
                                                                                                                                                                                                                                                                                              Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                        1 Similarity 100.0%; Score 21; DB 1; Le Similarity 100.0%; Pred. No. 6.1e+02; 1; Conservative 0; Mismatches 0;
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                                                                                                                                                                              Example 1; SEQ ID NO 41; 62pp; English
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                                                      Schroth GP;
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                                                                                                                                                                                                                                                                  artificial gene transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM07216 standard; DNA; 21
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                                                 Schroeder BG, Chen C,
(SCHR/) SCHROEDER B G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHEN/) CHEN C.
(SCHR/) SCHROEDER B.
(BRAN/) BRANDIS J.
(SCHR/) SCHROTH G.
                          SCHROTH G P.
                                                                                WPI; 2004-121562/12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      random primers.
                                                                                                                                                     polynucleotides.
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Best Local Simil
Matches 21; (
              CHEN
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                          (SCHR/)
              CHEN/)
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The present invention relates to a method for synthesising double-

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mixture comprising reverse transcriptase, RNA template, and first strain declared comprising reverse transcriptase, RNA template, and first strain declared council extranded cDNAs in a second reaction mixture comprising processive couple. Extranded cDNAs ligase, first CDNA strain as template and random to primers having a mixture of oligonucleotides having random DNA sequences. Also disclosed is a method for amplifying a population of RNA molecules to produce a pool of double-stranded cDNA molecules, and a kit for synthesising double-stranded cDNA. The generated cDNA products are useful in differential quantitative information about the genetic profile of nucleic acid in original RNA sample. The method of the invention is useful in differential gene expression assays for the analysis of diseased and normal tissue and for large-scale correlation studies on sequences, mutations, variants or polymorphisms among samples. The method is efficient in synthesising improved cDNA molecules and effective in generating useful quantities of an amplified CRNA product that comprises a population of CRNA molecules in substantially the same relative molar ratio as the RNA or mRNA starting material. The present sequence stranded cDNA, by synthesising first cDNA strands in a first reaction Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other; 

Gaps ö 0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; 0; Indels 100.0%; Pred. No. 6.1 tive 0; Mismatches 2709 AAAAAAAAAAAAAAAAA 2729 Query Match 0.8 Best Local Similarity 100. Matches 21; Conservative ઠે

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21 AAAAAAAAAAAAAAAAAA 1 硆

Allergic response suppressor oligonucleotide #912. ADU90228 standard; DNA; 21 BP 10-FEB-2005 (first entry) ADU90228; RESULT 368 ADU90228 

ss; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.

Synthetic.

US2004235774-A1.

25-NOV-2004.

23-APR-2004; 2004US-00831778

03-FEB-2000; 2000US-0179991P. 02-FEB-2001; 2001US-00776479.

(BRAT/) BRATZLER R L. (PETE/) PETERSEN D M. FOURON Y. (PETE/) (FOUR/) Petersen DM, Fouron Y; Bratzler RL,

WPI; 2004-833006/82.

Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.

Disclosure; SEQ ID NO 912; 235pp; English.

The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an

The invention relates to a novel glycosyltransferase capable of transferring an N-acetyl-D-galactosamine (GalNAC) residue to a GalNAC transferring an N-acetyl-D-galactosamine (GalNAC) residue to a GalNAC creceptor substrate The glycosyltransferase comprises a polypeptide having sequence ADV94808 containing amino acids of 43-601 or 1-601 of a fully defined sequence of 601 amino acids are in the specification, or ADV94808 in which one or more amino acids are comprises: a nucleic acid encoding the 601 amino acid glycosyltransferase protein and comprising a sequence ADV94807 having bases 127-1806 or 1-comprises: a nucleic acid encoding the 601 amino acid glycosyltransferase percification, or a sequence of 1806 base pairs, as given in the specification, or a sequence complementary to the 1806 bp that consists of the base sequence complementary to the 1806 bp complementary sequence; a recomblementary to the 1806 bp cor its complementary sequence; a recomblementary to the 1806 bp cor its complementary sequence; a recomblementary to the 1806 bp cor its complementary sequence; a recombinant containing the vector; an antibody capable of specifically recognizing the glycosyltransferase protein; and a protein; and a ö immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and acopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the method of the invention. GalNAc Novel glycosyltransferase capable of transferring N-acetyl-D-galactosamine (GalNAC) residue to GalNAC receptor substrate from GalNAC donor substrate, useful in screening substances that promotes/inhibits glycosyltransferase activity. glycosyltransferase; N-acetyl-D-galactosamine; GalNAc; screening; 88; Gaps ö 0.8%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 6.1e+02; rative 0; Mismatches 0; Indels Human glycosyltransferase pENTR/DTOPO vector 3' primer. Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other; (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO. (SEGK ) SEIKAGAKU KOGYO CO LTD. Example 1; SEQ ID NO 6; 37pp; Japanese. 2709 AAAAAAAAAAAAAAAAA 2729 21 AAAAAAAAAAAAAAAAAAA 06-JUN-2003; 2003JP-00162685 06-JUN-2003; 2003JP-00162685. ADV94812 standard; DNA; 21 10-MAR-2005 (first entry) Matches 21; Conservative WPI; 2005-035730/04. Local Similarity JP2004357635-A. 24-DEC-2004. PCR; primer. Synthetic. ADV94812; Query Match ADV94812/ 88888888888 ò В

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Gaps

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0; Indels

100.0%; Pred. No. 6.1e+02; ive 0; Mismatches 0;

21; Conservative

Best Local Similarity

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therapeutic agent of the disease caused due to change of activity of the glycosyltransferase, containing an active regulator of the glycosyltransferase protein as an active ingredient. The glycosyltransferase protein is useful in screening substances that promote or inhibit the activity of glycosyltransferase. The glycosyltransferase complementary DNA is useful as a probe for detecting in vivo expression of the glycosyltransferase protein is useful as the therapeutic of the diagnostic for medical studies. The active regulator of the glycosyltransferase protein, is useful as the therapeutic agent for treating the disease caused due to change of activity of the glycosyltransferase protein. The glycosyltransferase protein is capable for transfering dalNAc residue to a GalNAc receptor substrate from a GalNAc donor substrate. This polymuclectide sequence represents a primer used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel oligonucleotide molecule labeled with several fluorophores, useful for detecting biological molecules e.g., antibody, antigen, avidin or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 useful for detecting biological molecules e.g., antibody, antigen, avidin, protein, peptide, bacteria, virus, blood cell or hormone. ON is capable of providing strong fluorescence signals at different wavelengths. This sequence corresponds to an example of an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fluorophore-labeled biological detection oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                    0.8%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 6.1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADV86473 standard; DNA; 21 BP.
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                                                                                                                                                                                                                                                                                                                                                         Similarity
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Best Local
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0.8%; Score 21; DB 1; Length 21;

Query Match

Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;

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The invention relates to a complex (CI) comprising a first and second single stranded nucleic acid or its derivative and a metal ion, where the first and second base of the strands forms a bond with metal ion, where the included are detecting the existence of thymine-thymine, cytosine. Also circled are detecting the existence of thymine-thymine, cytosine cytosine-thymine base pair in a DNA or its analog (involving comparing DNA or its analog in an aqueous medium, processing the solution for 3 minutes, to obtain three DNA solutions, dissolving Hg(II)2+, Ag+ and combinations of Hg(II)2+ and Ag+ in the prepared DNA solutions, and comparing the existence of T-T base pair, C-C base pair and C-T base comparing the existence of T-T base pair, C-C base pair and C-T base comparing the existence of T-T base pair, C-C base pair and C-T base comparing the characteristic of metal binding region, where the coupling of metal ion (comprising one or more DNA molecules or their analogs having a metal binding region, where the coupling of metal ion is detected by analyzing the characteristic change in DNA). The complex (CI) is useful as a non-Watson Crick base pair in a double stranded DNA. The complex (CI) enables to detect non-Watson Crick base pair in a double stranded DNA. The complex (CI) enables to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Complex useful for detecting non-Watson Crick base pair in double stranded DNA, comprises first and second single stranded nucleic acid or its derivative and metal ion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence is a 21mer double stranded oligonucleotide with no Non-Watson-
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Pred. No. 6.1e+02;
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2709 AAAAAAAAAAAAAAAAA 2729
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                                                                                                                                BP
                                                                                                                                                                                                                                          Oligonucleotide DS21mer(A-T).
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26-FEB-2004; 2004JP-00051320.
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                                                                                                                            ADW71577 standard; DNA; 21
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Best Local Similarity
                                                                                                                                                                                                                                                                               DNA detection; ds
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                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                   ADW71577;
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ngs.res

Human KU70 transcript siRNA antisense oligonucleotide siRNA2.

(first entry)

28-JUL-2005

ADZ98948;

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The invention relates to interfering with transcription at a target site in a latent virus in a host cell. The method comprises producing an coligonucleotide capable of forming a triplex with the target site, coupling the oligonucleotide to a compound (A) capable of introducing transcription relations in the viral genome, introducing the coupled oligonucleotide, and reducing viral transcription from the target site. The method is used for interfering with transcription at target site in a latent virus, e.g. herpes viruses, UC virus, adenovirus, papillomavirus and HIV, and poxviruses, e.g. cowpox, monkeypox, camelpox, vaccinia, and variola in a host cell, and for effective anti-latency therapy to prevent recurrences and halt progression of harpes keratitis or genital herpes. The method enables control and prevention of infection through molecular transcriptional inhibition. The current sequence cepresents a triplex oligo (TFO) that binds target sequences in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Interfering transcription at target site in latent virus in host cell for treating viral infections comprises introducing triplex-forming oligonucleotide coupled to compound that introduces transcriptionaltering mutations in viral genome.
                                                                                                                                      Virucide; anti-HIV; viral transcription; inhibitor; infection;
herpes keratitis; genital herpes; ds.
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                                                                                                       Variola DNA binding triplex oligo (TFO), seq id 7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 7; 11pp; English
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ADY26140 standard; DNA; 21 BP.
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                                                                    05-MAY-2005 (first entry)
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JONES B B.
GRISSOM C B.
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                                                                                                                                                                                             Variola virus.
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                                                                                                                                                                                                                                                                                                                                                                                         (JONE/)
(GRIS/)
(HERP/)
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The invention relates to a method for modulating, in a host cell, a protein protein interaction between a first protein which is PRAK (P189-cepulated/activated protein kinase or MARKARKS) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host protein-protein interaction between a first protein which is PRAK and a second protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, quositis, Crohn's disease, gastritis, colitis, culcrative colitis, inflammatory bowel disease, proctitis, pelvic inflammatory disease, systemic lupus erythematosus, rhinitis, conjunctivitis, dermitis, dermitis or eczema. The present sequence represents an siRNA (short inferfering RNA) oligonucleotide targeting the KU70 transcript, which is used in the exemplification of the present
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protein interaction; short interfering RNA; siRNA; RNA interference;
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/*tag= a
/note= "2'-deoxythymine overhang"
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14-DEC-2000; 2000US-0255063P.
21-DEC-2000; 2000US-0256986P.
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15-MAR-2001; 2001US-0276179P.
19-MAR-2001; 2001US-0277013P.
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21-DEC-2001; 2001US-00024599.
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                                               gene silencing; ds.
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25-FEB-2000;
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Sequence 21 BP; 5 A; 7 C; 2 G; 2 T; 5 U; 0 Other;

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Gaps ö

Length 21; 0; Indels

Score 21; DB 1; Le Pred. No. 6.1e+02;

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ADZ98948 standard; RNA; 21 BP

RESULT 373 ADZ98948/ ID ADZ9

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cell a protein-protein interaction between a first protein which is PRAK
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                          Gaps
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        Length 21;
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      DB 1; Le
6.1e+02;
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/note= "2'-deoxythymine overhang"
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     0.8%; Score 21; DB 100.0%; Pred. No. 6.1 tive 0; Mismatches
                                          2036 AAGATGAAGGCTATCGTTGAG 2056
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                                                       AAGATGAAGGCTATCGTTGAG 1
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ADZ98946 standard; RNA; 21 BP.
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2000US-0256986P.
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2002US-00100503
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   Query Match 0.8
Best Local Similarity 100.
Matches 21, Conservative
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21-DEC-2001;
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23-JUL-2001;
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and a second protein which is BRK3 for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, juvenile chronic arthritis, myosilis, Crohn's disease, gastritis, colitis, ulcerative colitis, inflammatory bowelvesses, proteitis pelvic inflammatory disease, systemic lupus erythematosus, rhinitis, conjunctivitis, soleritis, chronic inflammatory polyneuropathy. Tertiary Lyme disease, psoriasis, dermatitis or eczema. The present sequence represents an siRNA (short interfering RNA) oligonucleotide targeting the furnity transcript, which is used in the exemplification of the present
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100.0%; Pred. No. 6.1e+02;
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/note= "2'-deoxythymine overhang"
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25-FEB-2000; 2000US-0185056P.
01-DEC-2000; 2000US-0025063P.
21-DEC-2000; 2000US-0255063P.
21-DEC-2000; 2000US-0256986P.
04-JAN-2001; 2001US-0259571P.
04-JAN-2001; 2001US-0259572P.
15-MAR-2001; 2001US-0276179P.
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21-DEC-2001; 2001US-00024599.
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The invention relates to a method for modulating, in a host cell, a protein-protein interaction between a first protein which is PRAK (P38-crequiated/activated protein kinase or MAPKARK5) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host cell a protein-protein interaction between a first protein which is BRAK for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, upositis, chronic arthritis, wyositis, Crohn's disease, proctiis, pelvic inflammatory disease, systemic lupus erythematosus, rhinitis, conjunctivitis, soleritis, chronic inflammatory disease, systemic lupus erythematosus, rhinitis, conjunctivitis, soleritis, chronic inflammatory polyneuropathy, Tertiary conjunctivitis, soleritis, chronic inflammatory polyneuropathy, Tertiary changes as psoriasis, dermatitis or eczema. The present sequence represents an sirNak (short interfering RNA) oligonucleotide targeing the KU70 transcript, which is used in the exemplification of the present
          Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPKS) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
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ö present invention relates to a method for augmenting T-helper 1 cells Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; immune stinulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colltis; eczema; skin allergy; contact dermatitis; ss; phosphorothioate. (Thi)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (1) to induce Thi-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (1) is useful for treating non-allergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease. comprises using a carbodismide derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid. (M1) is useful for labeling double stranded or single stranded nucleic acid or detecting single nucleotide polymorphisms. The present sequence was used to illustrate the method of the invention. Gaps ö Length 21; 0; Indels /note= "Phosphorothioate backbone" Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other; Score 21; DB 1; Lk Pred. No. 6.1e+02; 0; Mismatches 0; Immunostimulatory oligonucleotide, SEQ ID 881. Disclosure; SEQ ID NO 881; 58pp; English. contact dermatitis or latex dermatitis. 2709 AAAAAAAAAAAAAAAAAA 2729 Location/Qualifiers 0.8%; Sco. 100.0%; Prec 21 AAAAAAAAAAAAAAAAA 1 /\*tag= a /mod\_base= OTHER 29-MAR-2001; 2001US-0279642P. 29-MAR-2002; 2002US-00112653. 12-MAY-2005; 2005US-00127654 (IOWA ) UNIV IOWA RES FOUND AED75672 standard; DNA; 21 (first entry) Best Local Similarity 100. Matches 21; Conservative .. .21 \*tag= WPI; 2005-768014/78. Krieg AM, Berg DJ; US2005250726-A1 Key modified\_base 12-JAN-2006 10-NOV-2005 Synthetic AED75672; Query Match RESULT 377 AED75672/c 88888888888 ð 셤

Sequence 21 BP; 21 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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The new invention relates to preparing DNA fragments comprising sequences corresponding to two opposite end regions of a linear nucleic acid molecule, by creating a linear DNA molecule from a nucleic acid molecule, conference to two opposite ends of the linear DNA molecule, digesting the DNA molecule, circularizing the linear DNA molecule, digesting the circular DNA molecule, and endonuclease, and isolating the DNA fragment. Also described are a vector pGSC; obtaining (M2) information on the end sequences of a linear nucleic acid, comprising preparing DNA fragments by (M1), preparing a concatemer by ligating the DNA fragments by cother, and sequences of the linear nucleic; and priming (M3) a reverse cother, and sequences of the linear nucleic; and priming (M3) a reverse cother searched overhanging region, where the single-stranded overhanging region, where the single-stranded overhanging region, where the single-stranded overhanging region, to the single-stranded overhanging region, to the complementary 3 end sequence of the RNA; and letting the free 3'-end of the overhanging region of the linker prime a reverse transcription reaction over the RNA with a creverse transcription and letting the free 3'-end of the overhanging region of the linker prime a reverse transcription and expression profiling and for studies or biological system, characterization of genetic elements and analysis the expressed genes. The identified DNA fragments are useful in drug development, diagnostics or forensis ctudies. The present sequence is and proly. A prail removal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           opposite end regions of linear nucleic acid, for e.g. analysis, comprises ligating linkers, circularizing, and digesting.
                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA sequencing; gene expression; drug discovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Preparing DNA fragments comprising sequences corresponding to two
                                                                                                    ;
                                                          Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                        2709 AAAAAAAAAAAAAAAAAA 2729
                                                                                                                                                                                21 AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Fig 1, 70pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        priming and poly-A tail removal
                                                                                                                                                                                                                                                                                AEF40261 standard; cDNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JUL-2004; 2004WO-JP009862.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-JUL-2004; 2004WO-JP009862
                                                                                                                                                                                                                                                                                                                                                              23-MAR-2006 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosis; forensic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                        Poly A DNA sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Harbers M, Shibata Y;
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                                                        Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DNAF-) DNAFORM KK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-JAN-2006.
                                                                                                                                                                                                                                                                                                                          AEF40261;
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                                                                                                                                                                                                                                          Human interleukin-6 gene, herpes simplex, AIDS, modified, HIV, RSV, HPV,
malignancy, hepatitis, inflammation; ss.
                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.
                          ö
                                                                                                                                                                                                                    Oligomer IL6805 for forming triplex with HUMIL6 target duplex.
Query Match 0.8%; Score 21; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 6.1e+02; Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                         /*tag= d
/note= "o-xyloso dimer synthon linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= b
/mod_base= OTHER
/note= "OTHER= N4 N4 ethanocytosine"
                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod base= OTHER
/note= "OTHER= N4 N4 ethanocytosine"
ll. .12
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                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= c
/label= inverted_polarity_region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                     note= "see comments"
                                               2709 AAAAAAAAAAAAAAAAA 2729
                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                       1 AAAAAAAAAAAAAAAA 21
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                                                                                                                                 ВÞ
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91US-00686546.
91US-00686547.
91US-00766733.
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91US-00683420
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                                                                                                                                AAQ30432 standard; DNA; 23
                                                                                                                                                                                (revised)
(first entry)
                       21; Conservative
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                                                                                                                                                                                                                                                                                                               modified base
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07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                misc_feature
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17-APR-1991;
17-APR-1991;
27-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-NOV-1990
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08-APR-1991
                                                                                                                                                                                                                                                                             Synthetic.
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                                                                                                          RESULT 379
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interleukin 6 gene untranslated sequence contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HER, HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. The oligomer contains an inverted polarity region corned from an o-xyloso dimer synthon. The linking gp. is o-xyloso (nucleotides have the 3 positions of xylose sugars linked via the o-xylose ting). Two nucleotides are coupled through a xylene residue to contain the dimer synthon. This additional modifications may render the oligomer stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
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                       Сарв
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0
0.8%; Score 21; DB 1; Length 23; 00.0%; Pred. No. 6.4e+02;
                     0; Indels
       100.0%; Prec. ...
                       Matches 21, Conservative
             Best Local Similarity
  Query Match
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2709 AAAAAAAAAAAAAAAAA 2729 

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AAA29753 standard; DNA; 23 BP 15-AUG-2000 AAA29753; AAA29753/ 

Synthetic oligonucleotide #1. (first entry)

Primer; destabilise non-specific duplex formation; PCR; detection; purification; sequencing; genetic marker; RACE; DNA synthesis; ss.

Synthetic.

Location/Qualifiers /\*tag= a /mod\_base= i /note= "inosine" /note= "inosine" /\*tag= b /mod base= i Key modified\_base modified base

WO200020630-A1

13-APR-2000

99WO-CA000933 06-OCT-1999;

98CA-02246623

07-OCT-1998;

(UYMC-) UNIV MCGILL.

рав М; Pelletier J, WPI; 2000-328943/28.

Novel method of stabilizing duplex formation, or destabilizing non-specific duplex formation using primer containing modified nucleotide analogs, useful for preventing mispriming during PCR, RACE, DNA synthesis or sequencing

Example 1; Page 25; 46pp; English

The present invention describes a method for destabilising non-specific

The invention relates to systems for identification and characterisation

of gene expression in one or more samples, comprising an identimer having

a specific oligo-dT primer sequence, where the identimer comprises a

catectable marker at its 5' end. The system is useful for identifying any

cat all genes expressed in a given in vivo or in vitro RNA sample, as well

cas the relative differences in mRNA between 2 or more samples, where

cate expressed at different levels between 2 or more samples. The new

cystem or method addresses limitations of prior methods by comprising

compositions and systems that incorporate new strategies where molecular

compositions and systems that incorporate new strategies where molecular

compositions and systems that incorporate electrons are linked to DNA or RNA

sequence databases for optimal resource efficiency in assaying gene

expression. The system has the following advantages over existing

methods: (a) prior sequence information or clone library construction is ö duplex formation, between an oligonucleotide and a target nucleic acid (NA), comprising incubating the target NA with a modified oligonucleotide (I) comprising a homopolymeric sequence having a modification which decreases or abrogates H-bonding between the modified oligonucleotide and the non-specific target NA. The modified oligonucleotide is used to improve discrimination between the targeted homopolymeric sequence and a non-homopolymeric target sequence. It is used to increase the proportion of full length cDNA clones for a library, to reduce mispriming during sequencing, 5, or 3, RACE (rapid amplification of CDNA ends) or DNA synthesis or to generate bona fide genetic markers. The present sequence represents an oligonucleotide which is used in the exemplification of the Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identimer having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end. Gaps ö Oligo dT primer #2 used in method to study gene expression. Query Match 0.8%; Score 21; DB 1; Length 23; Best Local Similarity 91.3%; Pred. No. 6.4e+02; Matches 2; Indels Matches 2; Indels Oligo dT primer; gene expression analysis; primer; ss. Sequence 23 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 2 Other; 2709 AAAAAAAAAAAAAAAAAAAA 2731 Nagel AC; Disclosure; Page 11; 45pp; English. 23 AAAAANAAAAAAAAAAAAA 1 (GENO-) GENOMIC SOLUTIONS INC. В. 01-NOV-2000; 2000US-0244933P. 01-NOV-2001; 2001WO-US045401 ABK86169 standard; DNA; 24 24-SEP-2002 (first entry) Kane MD, Dombkowski AA, WPI; 2002-508123/54. present invention WO200236828-A2. 10-MAY-2002. Synthetic. ABK86169; RESULT 38 ABK86169 88888888888888888888888888 g ઠ

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The invention relates to systems for identification and characterisation of gene expression in one or more samples, comprising an identimer having a specific oligo-dT primer sequence, where the identimer comprises a catectable marker at its 5' end. The system is useful for identifying any or all genes expressed in a given in vivo or in vitro RNA sample, as well as the relative differences in mRNA between 2 or more samples, where the desired, for supporting discovery of new genes, and for identifying mRNAs that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising compositions and systems that incorporate new extrategles where molecular or biochemical assay compositions and systems are linked to DNA or RNA sequence databases for optimal resource efficiency in assaying gene expression. The system has the following advantages over existing methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence information in addition to information concerning changes or differences
                                                                                                                                                                                                                                                                                                         ö
not needed to enable the assay; (b) provides immediate sequence information in addition to information concerning changes or differences in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates CDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology the chinques; and (d) does not require prior knowledge of the sequence of the genome of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present sequence represents an oligo dT primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identimer having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.
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                                                                                                                                                                                                                                                                  Length 24;
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                                                                                                                                                                                                                         Sequence 24 BP; 20 A; 0 C; 1 G; 3 T; 0 U; 0 Other;
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Pred. No. 6.5e+02;
                                                                                                                                                                                                                                                                                                       Mismatches
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100.0%; Pre
0; 7
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ABK86168 standard; DNA; 24 BP.
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                                                                                                                                                                                                                                                                                                     21; Conservative
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                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-SEP-2002
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                                                                                                                                                                                   invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 382
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in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates CDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology techniques; and (d) does not require prior knowledge of the sequence of the ganome of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present sequence an oligo dT primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene using cells which contain defective mismatch generate genetically altered mutations with new
                                                                                                                                                                                                                                                                                                                                                               Hypermutable organism; dominant negative allele; mismatch repair gene; spontaneous mutation; DNA repair; purine nucleotide phosphorylase; PNP;
                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The patent discloses a method for generating hypermutable organisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kinzler KW
                                                                                                                                                                                                                                                                                                                                        Bacterial PNP DNA fragment with an out-of-frame polyA tract.
                                                                                                                                   Length 24;
                                                                                                                                                         0; Indels
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                                                                                                          Sequence 24 BP; 3 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
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"Out-of-frame polyA tract"
                                                                                                                                Score 21; DB 1; Le
Pred. No. 6.5e+02;
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100.0%; Pred. No. e...
'.. 0; Mismatches
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                                                                                                                                                                                                                                                                   BP
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VOGELSTEIN B.
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NICOLAIDES N
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                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200188192-A2
                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria.
Unidentified.
Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                         bacteria; ss.
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                                                                                    invention
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Best Local S
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(KINZ/)
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(MORP-)
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Length 26;

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Generating a mutation in a gene using a dominant negative allele of a mismatch repair gene which results in mismatch repair deficiency in cells containing the allele is useful in gene and drug target discovery and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to methods for generating a mutation in a gene of interesting using a dominant negative allele of a mismatch repair gene (I-MMR) under control of an inducible transcriptional regulatory element (ITRE). The invention is useful to provide new cell lines that can be used for gene discovery, drug target discovery, recombinant gene mutagenesis or recombinant protein production. The present sequence is a polyPNP (purine phosphorylase) out-of-frame polyA tract DNA
Dominant negative alleles of human mismatch repair genes can be used to generate hypermutable cells and organisms. They increase the rate of spontaneous mutations by reducing the effectiveness of DNA repair and thereby render the cells or animals hypermutable. The method is used to produce genetically altered organisms to produce new output traits. The present sequence is a bacterial poly purine nucleotide phosphorylase (polyPNP) DNA fragment containing an out-of-frame polyA tract. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dominant negative allele; mismatch repair gene; D-MMR; gene discovery; ITRE; inducible transcriptional regulatory element; recombinant gene mutagenesis; recombinant protein production; drug target discovery; ds.
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                                                                                                                                                                                      0.8%; Score 21; DB 1; Length 26; 00.0%; Pred. No. 6.7e+02; ve 0; Mismatches 0; Indels
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                                                                                                                                                      Sequence 26 BP; 22 A; 1 C; 2 G; 1 T; 0 U; 0 Other;
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                  AAD39650 standard; DNA; 26
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                                                                                                                                                                                                         Local Similarity 100.
1es 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nicolaides NC, Sass PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      recombinant technology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NICO/) NICOLAIDES N C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SASS P M.
GRASSO L.
VOGELSTEIN B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KINZLER K W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-499469/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2002055106-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                 AAD39650;
                                                                                                                                                                                           Query Match
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(GRAS/)
(VOGE/)
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                                                                                                                                                                                                                                                                                                                                                RESULT 384
                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                            SNP detection; breast tumor; endocrine disease; gynecology and obstetrics; neoplasm; cytostatic; metastasis; gene therapy; RNA interference; intercellular adhesion molecule; ICAM1;
                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying risk of, preventing and/or treating breast cancer by identifying and/or analyzing polymorphic variations in nucleotide sequences within the human genome.
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                                                                                                                                                                                                                                                                  Extend primer 57 used to genotype human ICAM region SNP DNA.
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0.8%; Score 21; DB 1; Length 26;
Best Local Similarity 100.0%; Pred. No. 6.7e+02;
Matches 21; Conservative 0; Mismatches 0; Indels
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 DB 1; Le
6.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kammerer SM,
0.8%; Score 21; DB
100.0%; Pred. No. 6.7
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 106; 617pp; English.
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                                                               2709 AAAAAAAAAAAAAAAAAA
                                                                                              AAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                  ICAM4; ICAM5; 88; PCR; primer.
                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUL-2003; 2003US-0490234P.
25-NOV-2003; 2003US-00723681.
25-NOV-2003; 2003US-0525239P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAY-2004; 2004WO-US016939.
                                                                                                                                                                         56
                                                                                                                                                                                                                                       (first entry)
Query Match 0.8
Best Local Similarity 100.
Matches 21; Conservative
                                                                                                                                                                         ADX99060 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2005-163257/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hoyal-Wrightson CR;
                                                                                                                                                                                                                                                                                                                                                                                                             WO2005014846-A2.
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                       05-MAY-2005
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                                                                                                                                                                                                         ADX99060;
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                                                                                                                                            RESULT 385
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RESULT 386 AAH24266/c

Sequence 26 BP; 22 A; 1 C; 2 G; 1 T; 0 U; 0 Other;

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The invention relates to human gonadotropin-releasing hormone 10 (AAM49158) and to nucleic acids encoding it (ABL55128). The protein has a molecular weight of 10 kb. The invention also relates to a method for the recombinant production of the protein, an antagonist of the protein, and the use of the protein, gene and antegonist in therapeutic applications. Gonadotropin-releasing hormone 10 can be used in the treatment of a variety of diseases such as cancer and HIV (human immunodeficiency virus) infection. Sequences ABL55129-ABL55130 represent reverse transcription-pck (RT-PCR) primers used in an exemplification of the invention to isolate human gonadotropin-releasing hormone 10 cDNA
    Human, gonadotropin-releasing hormone 10, recombinant production, cancer, HIV infection, human immunodeficiency virus, gene therapy, cytostatic, anti-HIV, reverse transcription-PCR, RT-PCR, primer, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polypeptide-human gonadotropin-releasing hormone 10 and polynucleotide encoding it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20.8; DB 1; Length 24; Pred. No. 6.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Extend primer 488 used to genotype human DPF3 SNP DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 1 A; 1 C; 3 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP detection; breast tumor; endocrine disease; gynecology and obstetrics; neoplasm; cytostatic; gene therapy; RNA interference; ss; PCR; primer; B4, zinc and double PHD fingers, family 3; DPF3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 17 (Disclosure); 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2704 GTACTAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                        (BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24 Grcccadadadadadadadada 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         guanine-nucleotide exchange factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%;
                                                                                                                                                                                                                                            31-MAY-2000; 2000CN-00116266.
                                                                                                                                                                                                                                                                                           31-MAY-2000; 2000CN-00116266.
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25-NOV-2003; 2003US-00723681.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-196660/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                         Xie Y;
                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                               CN1325900-A.
                                                                                                                                                                                              12-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                    Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
    8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to human phosphatase 79 (AAB73700), nucleic acids encoding it (AAH24264), and a method for the recombinant production of human phosphatase 79. The present invention additionally discloses an agonist of phosphatase 79 for therapeutic use, and an antibody which specifically binds to human phosphatase 79. Human phosphatase 79, and nuclectides which encode it may be used for treating a variety of nuclectides which encode it may be used for treating a variety of immunodeficiency virus) infection, immune disorders and inflammatory conditions. The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reaction or as a probe for hybridisation reactions, or in producing gene chips or microarrays. Sequences AAH24265-AAH24266 represent reverse transcription-per (1902) primers used in an exemplification of the invention to solate human phosphatase 79 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human phosphatase 79 and encoded polynucleotide, applicable in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunological diseases and various inflammation.
                                                                                                                                                                                 Phosphatase 79; human; BAC clone CTB-54D4-encoded protein homologue, recombinant production; malignant tumour; cancer; blood disease; HIV infection; human immunodeficiency virus; immunoder; inflammatory condition; oytostatic; anti-HIV; antiinflammatory; immunomodulator; reverse transcription-PCR; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human gonadotropin-releasing hormone 10 RT-PCR primer, SEQ ID NO:4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 2 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                       Human phosphatase 79 RT-PCR primer, SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BIOR-) BIOROAD GENE DEV LTD SHANGHAI.
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standard; DNA; 24 BP
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                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                               WO200138385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mao Y, Xie Y;
                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                           11-SEP-2001
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                                              AAH24266;
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metastasis;

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misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                       ABZ23535;
                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                          RESULT 390
                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                The invention relates to a novel method for identifying a subject at risk of breast cancer comprising detecting the presence or absence of a polymorphic variation associated with breast cancer. The method of the invention demonstrates cytostatic activity and may be useful for identifying a risk of, preventing and/or treating breast cancer and cancer metastasis. The methods may be utilized for gene therapy or RNA interference. The current sequence is that of an Extend primer of the invention which was used to genotype a human rho-family guaninenucleotide exchange factor D4, zinc and double PHD fingers, family 3 (DPF3) single nucleotide polymorphism (SNP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                               Identifying risk of, preventing and/or treating breast cancer by identifying and/or analyzing polymorphic variations in nucleotide sequences within the human genome.
                                                                                                                                                                                                                                                                                                                                           .;
0
                                                                                                                                                                                                                                                                                                                  / Match 0.8%; Score 20.8; DB 1; Length 26; Local Similarity 91.7%; Pred. No. 6.9e+02; nes 22; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunostimulatory non-CpG oligonucleotide IMT 173 SeqID 20.
                                            Reneland R;
                                                                                                                                                                                                                                                                                                Sequence 26 BP; 0 A; 3 C; 1 G; 22 T; 0 U; 0 Other;
                                            Kammerer SM,
                                                                                                                                                                                                                                                                                                                                                                 2709 AAAAAAAAAAAAAAAAAAAA 2732
                                                                                                                                               Example 16; Page 261; 617pp; English.
                                                                                                                                                                                                                                                                                                                                                                              Braun A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG75918 standard; DNA; 24 BP
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25-NOV-2003; 2003US-0525239P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IMMU-) IMMUNOTECH SA
                                           Roth RB, Nelson MR,
Hoyal-Wrightson CR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-053333/05.
                      (SEQU-) SEQUENOM INC
                                                                            WPI; 2005-163257/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG75918;
                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 389
ADG75918/c
                                                                                                                                                                                                                                                                                                                                             Matches
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This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CPG motif., which can stimulate an immune response in animals of the order of primate, including humans. The immune cresponse is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation, cytokine cytokine production in plasmacytoid dendritic cells. The present in plasmacytoid dendritic cells. The present in plasmacytoid dendritic cells. The present cinvention describes immunocodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinar DNA oligo in an ascendification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining whether a product of a gene is involved in preventing a replication error in a cell comprises providing a specific inhibitor for the product and determining the level of expression of a marker gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20.4; DB 1; Length 2
Pred. No. 7.1e+02;
0; Mismatches 1; Indels
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Claim 14; SEQ ID NO 20; 139pp; English.
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                          The specification describes a method for determining whether a product of a gene is involved in preventing a replication error in a cell. The method comprises providing the cell with a specific inhibitor for the product and determining the level of functional expression of a marker gene in the cell, where the level of expression of the marker gene is dependent on the occurrence of the replication error. The method is used for determining whether a product of a gene is involved in preventing a replication error in a cell. The identified genes are useful for developing diagnostic tools, or as targets for drug development to manipulate cells on the basis of the presence or absence of function of the gene. ABZ2335-36 represents fragments of plasmids used to detect somatic instability, in the course of the invention
                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for determining if a gene
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                                                                                                                                                                                                                                                            Score 20.4; DB 1; Length 25;
Pred. No. 7.2e+02;
0; Mismatches 3; Indels
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Example 1; Fig 3; 47pp; English
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22-MAY-2002; 2002WO-NLO00322.
28-NOV-2002; 2002WO-WO095071.
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/note= "N
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PLASTERK R H A.
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                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 21; Conserv
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product or compound is involved in preventing replication error in a cell. The method involves providing a cell with a specific inhibitor for a gene product or with a compound and determining the expression level of a marker gene in the cell, where the expression level of the marker gene is dependent on the occurrence of a replication error. The invention is useful in gene therapy and for treating a subject having tumours or cancer. The present sequence is a Caenorhabditis elegans heat-shock promoter DNA. This sequence is used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing a molecular array with a plurality of molecules immobilized to a solid substrate, useful in genetic analysis, gene expression studies or the detection or typing of single nucleotide polymorphisms in a sample of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention comprises a method for producing a molecular array, the method involves immobilising molecules to a solid phase at a density which allows individual immobilised molecules to be individually resolved. The molecular array produced by the method of the invention is useful for identifying one or more target molecules in a sample. The molecular array is also useful in genetic analysis, gene expression studies, identifying molecules which interact with a target molecule, detection/typing of single nucleotide polymorphisms, haplotyping and sequencing. The present DNA sequence represents a PCR primer that was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; target molecule identification; genetic analysis; SNP detection; haplotyping; sequencing; PCR; primer.
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                                                                                                                                                                                                                                                                3; Indels
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                                                                                                                                                                                        Sequence 25 BP; 21 A; 0 C; 1 G; 1 T; 0 U; 2 Other;
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Pred. No. 7.2e+02;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequencing. The present DNA sequence used in an example of the invention
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02-AUG-2001; 2001GB-00018879.
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21; Conservative
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AAL50570 standard; DNA; 22
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Best Local Similarity
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gene expression;
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                                                                                                                                                     invention.
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ACC48485
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mod_base= OTHER
note= "OTHER= locked nucleic acid"
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mod_base= OTHER
'not== "OTHER= locked nucleic acid"
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mod_base= OTHER
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mod_base= OTHER
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                                                                                          Locked nucleic acid anchored oligo(I) primer ON14.
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                                                                                                      Locked nucleic acid; LNA; gene therapy; primer;
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'mod_base= um
'note= "2'-0-methyluridine"
Mismatches
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            21 ВААААААААААААААААА
                                                       ACC48484 standard; DNA; 22 BP
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 Conservative
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 20;
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                                                                  ACC48484
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 Matches
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The present sequence is that of pyrene-anchored locked nucleic acid (LNA) coligo(dT) primer ON14, which was used in first-strand cDNA synthesis from enkaryotic mRNA. It includes compound '17d' at its 3' end, which is based on an LNA-type 2'-0,4'-C-methylene-beta-D-ribofuranosyl moiety. It is one of a set of such primers (see also ACC48482-85) that were used in an example from the invention to demonstrate improved reverse transcription of mRNA using pyrene-LNA anchored oligo(T) primers. The following results were observed: efficient priming on mRNAs with short poly(A) tails; curies resulting in an improved T20-VN anchor primer and thus avoiding reverse transcription of long poly(A) tracts; and improved reverse transcription of enkaryotic poly(A) tracts; and incorporated incorporated into nucleic acid strands. The novel LNA compounds can be used e.g. as into nucleic acid strands. The novel LNA compounds can be used e.g. as the primers, in sequencing, the synthesis of antisense oligonucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acid comprising a locked nucleic acid unit having a modified base that comprises an optionally substituted carbocyclic aryl moiety, or modified nucleobase or nucleosidic base other than oxazole/imidazole.
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Pred. No. 7e+02;
1; Mismatches 0; Indel8
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/mod_base= OTHER
/note= "OTHER= locked nucleic acid"
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/note= "OTHER= Compound 17d"
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22-SEP-2001; 2001US-0323967P.
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                                                                                                                                                                                                                                              04-SEP-2002; 2002WO-IB003911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 95.2
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kauppinen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-363021/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                            (EXIQ-) EXIGON AS
                                                                                    WO2003020739-A2
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-AUG-2003
                                                                                                                                                                     13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wengel J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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note = "OTHER = locked nucleic acid"

mod base= OTHER

modified base

/\*tag≂ a

/\*tag= b /mod\_base= OTHER /note= "OTHER= locked nucleic acid"

/\*tag= d /mod\_base= OTHER /note= "OTHER= locked nucleic acid"

'mod\_base= OTHER 'note= "OTHER= locked nucleic acid"

O

\*tag=

modified base

modified\_base

/\*tag= e
mod\_base= OTHER
note= "OTHER= locked nucleic acid"

/mod

modified base

modified base

modified base

/\*tag= g /mod\_base= OTHER /note= "OTHER= locked nucleic acid"

/note= "OTHER= Compound 17d"

WO2003020739-A2

13-MAR-2003.

04-SEP-2002; 2002WO-IB003911. 04-SEP-2001; 2001US-0317034P. 22-SEP-2001; 2001US-0323967P.

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Wengel J, Kauppinen

(EXIQ-) EXIQON AS.

base= OTHER

\*tag= h

modified\_base

'note= "OTHER= locked nucleic acid"

/\*tag= f /mod base= OTHER

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The present sequence is that of pyrene-anchored locked nucleic acid (LNA) cligo(dT) primer ON15, which was used in first-strand cDNA synthesis from eukaryotic mRNA. It includes compound '17d' at its 3' end, which is based on an LNA-type 2'-0,4'-C-methylene- beta-D-ribofuranosyl moiety. It is come of a set of such primers (see also ACG48482-84) that were used in an example from the invention to demonstrate improved reverse transcription of mRNA using pyrene-LNA anchored oligo(T) primers. The following results were observed: efficient priming on mRNAs with short poly(A) tails; efficient anchoring of the oligo(T) primer by pyrene-LNA and LNA-C/G/T con this resulting in an improved T2O-VN anchor primer and thus avoiding reverse transcription of eukaryotic poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; con continue avoiding that comprise unique base groups. Desirable nucleobase and nucleosidic chase substitutions can mediate universal hybridisation when incorporated into nucleic acid strands. The novel LNA compounds can be used e.g. as CRC primers, in sequencing, the synthesis of antisense oligonucleotides, and in Alamoneric
                                                                                                                                                                                                                                                                                                                                         Novel nucleic acid comprising a locked nucleic acid unit having a modified base that comprises an optionally substituted carbocyclic aryl molety, or modified nucleobase or nucleosidic base other than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;
                                   /mod_base= OTHER
/note= "OTHER= Compound 17d"
                                                                                                                                                                                                                                                                                                                                                                                                                               Example 24a; Page 90; 119pp; English.
                                                                                                                                                                                      04-SEP-2001; 2001US-0317034P. 22-SEP-2001; 2001US-0323967P.
                                                                                                                                                    04-SEP-2002; 2002WO-IB003911
               *tag= b
                                                                                                                                                                                                                                                                         Wengel J, Kauppinen S;
                                                                                                                                                                                                                                                                                                       WPI; 2003-363021/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in diagnostics
                                                                                                                                                                                                                                      (EXIQ-) EXIQON AS.
                                                                                                                                                                                                                                                                                                                                                                                              oxazole/imidazole.
                                                                                    WO2003020739-A2
modified_base
                                                                                                                    13-MAR-2003.
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                            Gaps
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0
  DB 1; Length 22;
                          0; Indels
0.7%; Score 20.2; DB 1
95.2%; Pred. No. 7e+02;
tive 1; Mismatches
                                                2708 TAAAAAAAAAAAAAAAAAAAA 2728
                                                               Best Local Similarity 95.2
Matches 20; Conservative
  Query Match
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Locked nucleic acid; LNA; gene therapy; primer; ss. Locked nucleic acid anchored oligo(I) primer ON13 ACC48483 standard; DNA; 22 BP. 11-AUG-2003 (first entry) ACC48483; ACC48483/ 14x8x4x8x6x4x6

Location/Qualifiers 2

Key modified\_base

Synthetic.

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"... recovers and the control of pyteune-aucunoted located (LNA) oligo(dT) primer ON13, which was used in first-strand cDNA synthesis from cukaryotic mRNA. It includes compound '17d' at its 3' end, which is based on an LNA-type 2' -0,4' -C-methylene- beta-D-ribofuranosyl moiety. It is cone of a set of such primers (see also ACC48482-85) that were used in an example from the invention to demonstrate improved reverse transcription of mRNA using pyrene-LNA anchored oligo(T) primers. The following results were observed; efficient priming on mRNAs with short poly(A) tails; cefficient anchoring of the oligo(T) primer by pyrene-LNA and LNA-C/G/T units resulting in an improved T2D-VN anchor primer and thus avoiding ceverse transcription of long poly(A) tracts, and improved reverse transcription of eukaryotic poly(A) tracts, and improved reverse transcription of elong poly(A) anchor primer and thus avoiding transcription of elong poly(A) and incectly from total RNA extracts due to increased specificity. The invention relates to modified LNA units that comprise unique base groups. Desirable nucleobase and nucleosidic into mucleic acid strands. The novel LNA compounds can be used e.g. as into midianostics that computing the synthesis of antisense oligonucleotides, and in diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of pyrene-anchored locked nucleic acid (LNA)
                                                                                                          Novel nucleic acid comprising a locked nucleic acid unit having a modified base that comprises an optionally substituted carbocyclic aryl moiety, or modified nucleobase or nucleosidic base other than oxazole/imidazole.
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WPI; 2003-363021/34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method for assessing a condition of a performance animal. The method involves determining in sample abundance of expressed target nucleic acid; transmitting digital sample signal to remote diagnostic server; processing digital sample signal to remote diagnostic server; processing digital sample signal at remotely located database to correlate digital signal with digital information and returning report of particular condition of animal. The method is useful for assessing a condition of a performance animal, and be an athletic ability and a condition that enhances, hinders, impedes or does not change an expected ability of the performance animal; and also normal, pre-clinical, overt progress and/or stage of disease, undiagnosed of unclassified conditions, presence of drugs, response to expriremental conditions. Diseases assessed by the states and response to environmental conditions. Diseases assessed by the invention include laminitis, lameless, viral or bacterial disease, gastrits, gastric ulcers, respiratory aliments, fractures, epistaxis, musculoskeletal damage or disorders and joint diseases. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states, of performance animal involves analyzing nucleic acid expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laminitis; viral disease; vaccine; bacterial disease; primer; epistaxis; gastritis; gastric ulcer; respiratory ailment; fracture; joint disease; musculoskeletal damage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anchored oligo dT primer used to illustrate the method of the invention.
                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence is a primer used to illustrate the method of the invention
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                                         0.7%; Score 20.2; DB 1; Length 22;
95.2%; Pred. No. 7e+02;
tive 1; Mismatches 0; Indels
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Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENO-) GENOMICS RES PARTNERS PTY LTD
                                                                                                                                           2708 TAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 46; 87pp; English.
                                                                                                                                                                                          BAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAY-2002; 2002WO-AU000553
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29-JUN-2001; 2001US-00896941
                                                                                                                                                                                                                                                                                                           AAD51324 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-2003 (first entry)
                                         Query Match
Best Local Similarity 95.27
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brandon RB;
                                                                                                                                                                                                                                                                                                                                                          AAD51324;
                                                                                                                                                                                                                                                               RESULT 396
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The invention relates to methods and system for sequence presentation. The method involves extracting a partial sequence corresponding to a partial sequence of a target gene having an unidentified sequence, by homology search on a database. The methods are useful for presentation of sequences. It is also useful to prepare primer sequences to obtain a cDNA clone of a total coding region from a partial sequence of a gene having an unidentified sequence. The present sequence is a primer specific for human chromosome 10 RP-11-336AlO clone DNA. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Presenting partial sequences by predicting and extracting exon sequences from a database, is useful to prepare primers to obtain a cDNA clone of total coding region from a partial sequence of an unidentified gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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95.2%; Pred. No. 7e+02;
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                                                                                                                                                                                                                                                                                                                                  Sequence presentation; human; chromosome 10; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Mismatches
                                                                                                                                                                                                                                                                                     Human RP-11-336A10 clone specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; SEQ ID NO 56; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2708 TAAAAAAAAAAAAAAAA 2728
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BP.
                                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-APR-2002; 2002JP-00103333.
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                                                                                                                                              22
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                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                              AAD64451 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hosoiri T, Yokoi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-864174/80.
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                                                                                                                                                                                                                                                                                                                                                                                                                               US2003190648-A1.
                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAR-2003
                                                                                                                                                                                                                                           12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-OCT-2003.
                                                                                                                                                                                             AAD64451;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABX74887;
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                           21
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ABX74887/c
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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a microarray comprising a matrix of at least one CDNA probe from a set of probes immobilised to a solid surface in a predetermined order, where a row of pixels corresponds to replicates of one distinct probe from the set. The probes are complementary to nucleic acid sequences that are expressed differentially in aggressive as compared to non-aggressive types of clear cell renal carcinoma (CC-RCC) and that hybridise to the probes under high stringency conditions. The microarray is useful for the prognosis of patients with CC-RCC, wherein aggressive and non-aggressive CC-RCC tumour types are characterised by differential expression profiles of genes that hybridise with one or more probes immobilised on the microarray. The arrays are useful for sene expression profiling of tumour and normal tissues. The present sequence represents an oligo-dT primer used in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                           New microarray, comprising a matrix of cDNA probe from a set of probes immobilized to a solid surface in predetermined order, useful in the prognosis of patients with clear cell renal carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer; neuroblastoma; rhabdomyosarcoma; Burkitt's tumour family;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
 gene expression profiling; tumour tissue; oligo-dT; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20.2; DB 1; Length 22;
Pred. No. 7e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA extraction anchored oligonucleotide primer.
                                                                                                                                                    Takashi M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2708 TAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                       Example 2; Page 30; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 BAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADI34007 standard; DNA; 22 BP
                                                                                    29-MAR-2002; 2002WO-US009576.
                                                                                                         29-MAR-2001; 2001:US-0279411P.
                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-APR-2002; 2002US-00133937
                                                                                                                                                    BT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ewing tumour family; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-APR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20; Conservative
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                                                                                                                            (VAND-) VAN ANDEL INST
                                                                                                                                                                       WPI; 2003-040679/03.
                                                                                                                                                   Rhodes D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS2004009154-A1
                                         WO200279411-A2
                                                              10-OCT-2002
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                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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The invention relates to a method of selecting genes expressed in a cancer cell, which involves characterising cancer based on the functioning of gene selection by comparing the expression of the selection of gene from the cancer cell with the expression of an identical selection of genes from a noncancerous cell or different type of cancer cell. The cathod is useful for targeting genes expressed in a cancer cell. The method is useful for targeting the therapy of cancer by using a selection of genes or their product sexpressed in a cancer cell, the gene selection of genes or their product sexpression of the selected gene or their products from the comparing the expression of the selected gene or their products enocancerous. The method is also useful for diagnosing, prognosing, monitoring and classifying a disease condition e.g. curous. The present sequence represents an anchored oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
0
                                                                                                                                                                                                                                        Selecting genes expressed in cancer cell, by characterizing cancer based on functioning of gene selection by comparing expression of selected gene from cancer cell with expression of selected genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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35.2%; Pred. No. 7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYCH-) UNIV CHANCELLOR MASTER & SCHOLLARS OXF.
                                                                                                                                          Meltzer P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 18; 53pp; English.
                                                                                                                                          Peterson C,
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26-SEP-2002; 2002GB-00022412.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; primer; molecular array.
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KHAN J.
RINGNER M.
PETERSON C.
MELTZER P.
                                                                                                                                       Khan J, Ringner M,
                                                                                                                                                                                         WPI; 2004-167702/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                 (RING/)
(PETE/)
     KHAN/)
                                                                                MELT/)
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ADL97794/c
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02-DEC-2004
                                                             Synthetic
                                            ADS13095;
                              Query Match
                                        RESULT 401
                                Matches
                                         ADS13095,
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The invention relates to a method of producing (MI) a molecular array, involves: immobilizing to a solid phase a several molecules at a density which allows individual immobilized molecules to be individually resolved, where each molecule in the array is spatially addressable and the identity of each molecule is known or determined prior to immobilization; and optionally providing a molecular array comprising a several molecules immobilized to a solid phase at a density such that individual immobilized molecules are not capable of being individual resolved, and reducing the density of functional immobilized molecules in the array such that remaining individual functional immobilized molecules in the resulting array is spatially addressable and functional molecule in the resulting array is spatially addressable and the identity of each molecule is known or determined prior to the density reduction seep. The array efficiently resolve complex samples, separate correct signals from erroneous signals, eliminates need for sample correct signals from erroneous signals, eliminates need for sample of single molecule processes. This sequence represents anoligonucleotide used in the method of the invention. ö Producing molecular array by immobilizing to solid phase several known molecules at low density for allowing individual immobilized molecules to be individually resolved and spatially addressable. in vitro evolution; hyperlipidaemia; atherosolerosis; neurological disease; Alzheimer's disease; neurofibromatosis; Huntington's disease; depression; amyorrophic lateral sclerosis; multiple sclerosis; stroke; Parkinson's disease; multi-infarct dementia; cancer; organophosphate poisoning; antilipaemic; antiarteriosclerotic; neuroprotective; nootropic; cytostatic; anticonvulsant; antidepressant; antiparkinsonian; antidote. paraoxonase; PON1; praoxon; nerve agent; sarin; soman; Gaps ö Oligo dT PCR primer used in the cloning of PON1 genes Seq 11. / Match 0.7%; Score 20.2; DB 1; Length 22; Local Similarity 95.2%; Pred. No. 7e+02; nes 20; Conservative 1; Mismatches 0; Indels Silman I; Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other; Sussman JL, Disclosure; Page 152; 219pp; English Gaydukov L, 2708 TAAAAAAAAAAAAAAA 2728 21 BAAAAAAAAAAAAAAAAA 1 (YEDA ) YEDA RES & DEV CO LTD ADS13095 standard; DNA; 22 BP 04-MAR-2004; 2004WO-IL000216. 04-MAR-2003; 2003US-0451267P. 22-OCT-2003; 2003US-0512925P. (first entry) Aharoni A, WPI; 2004-668627/65. primer; 88; WO2004078991-A2. 16-SEP-2004 rawfik DS,

Novel mutant of PON enzyme exhibiting increased substrate specificity to

This invention relates to novel mutant serum paraoxonase (PON1) nucleic acid molecules and the encoded proteins thereof. Specifically, it refers to enzymes that are calcium dependent phosphoricistcrases essential to the detoxification process of organophosphates such as the insecticide praoxon and the nerve agents sarin and soman. The present invention describes a method to identify mutated PONs that exhibit substrantially identical (or improved) substrate specificity in comparison with the wild crype PON and also those mutants that do not form aggregates when expressed in bacteria. In particular, the method employed an in vitro expressed in bacteria. In particular, the method employed an in vitro confuncing plasticity, catalytic activity and maintaining substrate binding. These mutants have been found to be useful for treating or preventing PON1-related diseases including hyperlipidaemia.

Confuncial plasticity, catalytic activity and maintaining substrate concupity mutiple scalerosis, mutiple scales including hyperlipidaemia.

Confuncial mutiple scalerosis, stroke, parkinson's disease, they exhibit antilipaemic, antiarteriosclerotic, neuroprotective, concurropic, cytostatic, anticonvulsant, antidepressant and case of conting and expression of a wild type PONI gene of the invention. PCR primer to amplify the 5' CDS of rice Xa31 cDNA by RACE PCR Seg 23b. New isolated nucleic acid sequence that provides resistance to plants, useful for making transgenic plants that are resistant to bacterial blight disease caused by Xanthomonas. PON substrate, useful for treating or preventing PON1-related diseases e.g., hyperlipidemia, atherosclerosis, neurological disease or cancer. Gaps plant, disease resistance, crop improvement, transfection, transgenic plant, bacterial blight disease; primer; ss; PCR; RACE ö 0.7%; Score 20.2; DB 1; Length 22; 95.2%; Pred. No. 7e+02; ive 1; Mismatches 0; Indels 7e+02; 0; Seguence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other; Example 1; SEQ ID NO 11; 240pp; English. 2708 TAAAAAAAAAAAAAAAAA 2728 ; K Example 7; Page 35; 93pp; English 21 BAAAAAAAAAAAAAAAAA 1 ច្ច BP. (TEMA-) TEMASEK LIFE SCI LAB. 13-AUG-2003; 2003WO-SG000191. 13-AUG-2003; 2003WO-SG000191. 95.2%; Wang G, Tian DS, ADY30380 standard; DNA; 22 (first entry) 20; Conservative WPI; 2005-182374/19. Best Local Similarity WO2005017158-A1 Oryza sativa. 05-MAY-2005 24-FEB-2005. ADY30380; Query Match Yin ZC, Matches ADY30380/ ##X#XDDDDDDDDDDDDDDDDDDDDXX 셤 ઠે

Page 195

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This invention relates to a novel nucleic acid molecule that provides a plant with resistance to Xanthomonas when transfected into that plant. Specifically, it refers to a method of generating a plant that is resistant to Xanthomonas by transfecting the resistance gene Xa31 into the plant or transfecting Xa31 into a plant cell (or cells) and growing a plant thereof. The present invention describes an appropriate vector comprising a plant promoter operably linked to the nucleic acid, which can be used to produce the transgenic plant, preferably transgenic rice. As such, Xa31 acid is useful for generating transgenic plants that are resistant to bacterial blight disease caused by the bacterium Xanthomonas. This oligonucleotide sequence is a PCR primer given in an exemplification of the invention. NOTE: This sequence taken from example 7 differs from that given in the sequence listing of the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for providing a profile of mRNA molecules present in a sample. The method comprises generating two independent patterns characteristic of the population of mRNA molecules expressed in the sample and analysing the patterns using a combinatorial algorithm, comparing gene expression by different or same cell types andorf different conditions, and identifying genes having a role in various cellular processes. The method is useful for the analysis and identification of transcribed genes, and fingerprinting. The method can be used to identify genes which play a role in determining various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Providing mRNA profile, by generating two independent patterns characteristic of sample mRNA population, analyzing patterns, comparing gene expression by cell types under varied conditions, and identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identification of transcribed gene; mRNA profile; gene expression; cellular process; fingerprinting; susceptibility to external factor; development; disease; PCR; primer; ss.

    PCR primer used in method of identifying transcribed genes.

                                                                                                                                                                                                                                                                                                                                                               Score 20.2; DB 1; Length 22;
Pred. No. 7e+02;
1; Mismatches 0; Indels
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21-JUL-2000; 2000US-0219925P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying extendible primers for use in identification, or classification of a nucleic acid of an organism, allele or gene such as class 1/2 HLA comprises identifying all possible nucleotide sequences of specific length.
cellular processes, including susceptibility to external factors, development, and disease. The present sequence for a PCR primer is used in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                             DNA sequence analysis; sequencing; protein sequence; protein structure; gene typing; organ donation; bacteria identification; 16s rRNA; HLA; human leukocyte antigen; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides a method for identifying a set of extendible primers which can be used in the identification, typing and classification of genes. This can then be used to predict protein sequence and structure, in organ donation to match the organ with the receiver, and to identify bacteria in a sample. The method can be used type the human leukocyte antigen genes (HLA) and 16s rRNA genes in
                                                                                                                            Gaps
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                                                                                            0.7%; Score 20.2; DB 1; Length 23; 95.2%; Pred. No. 7.1e+02;
                                                                                                                            0; Indels
                                                               Sequence 23 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 3 Other;
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                                                                                                           Pred. No. 7.1e
1; Mismatches
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                                                                                                                                                                                                                                                                   AAC96256 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                HLA DPA1 gene PCR primer #13.
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hes 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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Best Local S
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RESULT 405 ABA03917/c

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                                                                                                                                                                                                                                                                                                                                           The present invention describes human connexin 9 (1). (1) has cytostatic, virucidal, immunomodulatory, antiinflammatory and haemostatic activities. (1) and the polynucleotide encoding it (II) are applicable in the diagnosis and treatment of malignant tumour, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence represents a PCR primer for human connexin 9, which is used in an example from the present invention
                                                                                                                                                                                                                                                                               Human connexin 9 and encoded polynucleotide, applicable in diagnosis and treatment of malignant tumor, hemopathy, HIV infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow; microsatellite; PCR; primer; ss.
                                                                              Human; connexin 9; cytostatic; virucidal; immunomodulatory;
antiinflammatory; haemostatic; malignant tumour; haemopathy;
HIV infection; immunological disease; inflammation; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 3 A; 0 C; 1 G; 21 T; 0 U; 0 Other;
                                                           Human connexin 9 PCR primer 2 SEQ ID NO:4.
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  ABA03917 standard; DNA; 25 BP.
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                                                                                                                                                                                                     27-APR-2000; 2000CN-00115456
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                                         (first entry)
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                                                                                                                                           WO200181538-A2
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                                                                                                                         Homo sapiens
                                         18-FEB-2002
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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA sample containing at least one segment of (1); and determining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or compared pML; and prediagnosis (M3) of diseases associated with gene that include PML; The method is used to identify microsatellite markers, in a include PML. The method is used to identify microsatellite markers, in a diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperhermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, cometabolic diseases, also to type genes that encode milk proteins, particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion cycle protein (PrP) comprising a polymorphic microsatellite locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 20.2; DB 1; Length 25; 88.0%; Pred. No. 7.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 0 A; 3 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cow prion protein microsatellite locus primer #72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25 AAAAAAAAAAGAAAGAAAGAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 28; 64pp; German.
                                                                                                                                                                                                                                                                                           Han Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    microsatellite; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2002; 2002DE-01036711
                                                                      09-AUG-2002; 2002DE-01036711.
                                                                                                                                            09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-2002; 2002DE-01036711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADO81060 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22; Conservative
                                                                                                                                                                                                                     (UYHO-) UNIV HOHENHEIM
                                                                                                                                                                                                                                                                                               Geldermann H, Preuss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     length of amplicons.
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-215730/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DE10236711-A1.
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26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Typing
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                                                                                                                                                                           The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA semple containing at least one segment of (1); and determining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or more PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a include PML. The method is used to identify microsatellite markers, in a classase and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion cycle protein (PPP) comprising a polymorphic microsatellite locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                          genes that contain polymorphic microsatellite loci, useful for fying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 20.2; DB 1; Length 25; 88.0%; Pred. No. 7.4e+02; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA synthesis; RNA; antisense strands; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25 BP; 0 A; 3 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dye-coupled 3'-amino modified oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a ./note= "3-amino modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mag M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAAAAAAAA 2733
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Konrad R,
                                                                                                                                                     Example 3; Page 28; 64pp; German.
                                 Han Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ25565 standard, DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         91EP-00120935
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                              Geldermann H, Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 88.0
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Engels J, Herrlein M,
(UYHO-) UNIV HOHENHEIM
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02-DEC-1992 (first en
                                                                                                          identifying predispo
length of amplicons.
                                                            WPI; 2004-215730/21
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ25565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                          Pyping
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                                   New dye-coupled modified nucleosides, nucleotides and oligo:nucleotides useful for synthesis of antisense DNA and RNA strands in presence of template, also for in-vivo and in-vitro detection of genetic material.
                                                                                                                                                              The sequence is an example of a dye coupled 3'-amino modified oligonucleotide, it can be used in the synthesis of DNA and RNA nucleosides, nucleotides and oligonucleotides and for the synthesis of opposite strands in the presence of template strand and in fluorescence microscopic and macroscopic detection in vivo and in vitro of genetic material. It is labelled with a fluorescent dye. See also AAQ25566 and AAQ25567. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.88+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Microsatellite sequence from clone AGLA247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 AAAAAAAAAAAAAAAA 1
                                                                                                                         Example; Page 9; 17pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ33554 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92WO-US000340
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                        20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1992-284684/34.
WPI; 1992-201578/25,
                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ33554;
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(CROS/) CROS P.
(KURF/) KURFURST R.
(BATT/) BATTAIL N.
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                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                         10-JUN-1994;
                                                                                                                                                                                         misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-1993;
                                                                                                                                                                                                                                                                                                                   WO9429723-A1
                                                                                                                                                                                                                                                                                                                                                22-DEC-1994.
                                       25-MAR-2003
24-AUG-1995
                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
               AAQ94205;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The inventors claim a method of transporting a nucleic acid deriv. accros a membrane which comprises using a receptor that uses salt bridgin, aromatic setacking, H bonding and chelation to recognise the nucleic acid deriv. AAQ58305, AAQ58577-86 are nucleic acid derivs used in the examples. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transport of nucleic acid derivs. across membranes - using new receptors which use salt bridging, aromatic stacking, hydrogen bonding and
used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                            Sequence of synthetic RNA oligo which is a target nucleotide for a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                          Gaps
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100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                             Length 20;
                                                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                       Novel receptor; nucleic acid; transport; oligo; ss.
                                                                                  Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                        0
                                                                                                             DB 1; Le
6.8e+02;
                                                                                                          Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Table 1, page 38; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  De Mendoza
                                                                                                                                                                                                                                                                 AAQ58578 standard; RNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93WO-US007603
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                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-AUG-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-1992;
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                                                                                                                                                                                                                                                                                                                       25-MAR-2003
21-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chelation.
                                                                                                                                                                                                                                                                                                                                                                                receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Usman N,
                                                           field.)
                                                                                                                                                                                                                                         RESULT 410
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AAQ94205 standard; DNA; 20

AAQ94205/c ID AACO

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Assay device for hapten or its specific antibodies - comprises support having competitive reagent immobilised via nucleic acid ligand to improve orientation and accessibility.
                                                                                                                                                                                                                                                                                                                                     /*tag= b /nctag= b /nctag= nthe glycosidic bonds between nucleotides are all in the alpha-anomer form"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "carries a group derived ffrom aminopropanediol"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                            Alpha-anomeric oligonucleotide ligand 1803 for oestradiol hapten.
                                                                                                                                      Oligonucleotide ligand; steroid hormone; hapten; immobilisation; immunodetection; estradiol; alpha-anomer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Score 20; DB 1; Le
100.0%; Pred. No. 6.8e+02;
:ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 10; 39pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Battail N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AAAAAAAAAAAAAAA 1
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                              (first entry)
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Matches 20; Conservative
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*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=
(revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-036665/05.
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Tue Nov

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                                                                                                                                                                                                                                                                                                                                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                           Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T2; HLA; dQa; self-addressable electronic device; SAED; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/note= "3' aminolink2 Thymine; allows binding to any
amine"
                                                                    Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
0
                                        Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               T2 (synthetic DNA probe with 5' amino terminal #4)
                                                                                                                                                                                                                                                     (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2706 ACTAAAAAAAAAAAAAA 2725
                                                                                   restriction enzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 ACTAAAAAAAAAAAAAAA
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             04-AUG-1995 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 20; Conservative
                                                                                                                                                                                                                                                                                WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                       JP06303997-A
                                                                                                                                                                                            16-APR-1993;
                                                                                                                                                                                                                          16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9512808-A1
                                                                                                                                                                   01-NOV-1994
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                                                                                   aggregate;
                                                                                                             Synthetic.
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Best Local (
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The sequences represented by, AAQ90402-15 are synthetic DNA probes containing 5' amino termini. The sequences shown in AAQ90390-401 are synthetic DNA probes with 3' ribonucleoside termini. Thes sequences were specific for the polymorphisms of HLA gene dQa. The sequences were in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and sensitivity of these reactions are greatly improved at micro-locations
                                                                                                                        New self-addressable electronic devices - used for multi-step and multiplex reactions such as DNA hybridisation(s), clinical diagnostics and bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalian stem cell factor (SCF) cDNA oligonucleotide primer 220-7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stem cell factor; progenitor; haematopoiesis; SCF; anaemia; thrombocytopenia; leucopenia; AIDS; immunodeficiency; bone graft; transplant; neoplasia; myelosuppression; bone marrow; ss.
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                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                     Seguence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                   Example 1; Page 41; 86pp; English.
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90US-00537198.
90US-00573616.
90WO-US005548.
             93US-00146504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90US-00589701
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                                                                                                                                                                                                                                                                                                                                                                                                                             20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
                                                                                               WPI; 1995-185870/24.
                                       (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                  Heller MJ, Tu
             01-NOV-1993;
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15-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-OCT-1990;
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28-SEP-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 414
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New stem cell factor polypeptide(s) - for stimulating the growth of

Martin FH;

Bosselman RA,

Suggs SV,

Zsebo KM,

(AMGE-) AMGEN

WPI; 1995-346090/45

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AAT04915-T04922 are oligonucleotide primers and probes used for the amplification and sequencing of mammalian stem cell factor (SCF). Non-maturally occuring SCF and of-terminally truncated polypeptides, having amino acid sequences sufficiently duplicative of naturally ocurring SCF, stimulate growth of primitive progenitors such as haematopoietic progenitor cells, neural stem cells and primordial germ stem cells. The perides can be used in a composition for treating leucopenia, anaemia or thrombocytopenia, for enhancing engraftement of bone marrow during transplantation or for bone marrow recovery after chemotherapy or cadiation-induced bone marrow aplasia or myelosuppression. They can also be used for treating neoplasia, nerve damage, infertility, intestinal damage or myeloproliferative disorders. Antibodies may be raised against the peptides for use in detection or neutralisation of SCF in serum. SCF may be useful for the treatment of AIDS and severe combined immunodeficiency (SCID) states alone or in combination with other factors such as IL-7. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
primitive progenitor cells, esp. for treating disorders involving blood
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sulphurisation of phosphorus-containing compounds, e.g. oligo:nucleotide(s) - by contacting the compound with a dissulphide-containing five-membered heterocycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phosphorothioate; sulphurisation; heterocycle; automated synthesis; antisense; EDITH; Beaucage reagent; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/note= "phosphorothioate internucleotide linkages"
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0
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                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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(LOUU ) UNIV LOUISIANA STATE & AGRIC.
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                                                    Example 3; Fig 12C; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2707 CTAAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Phosphorothioate oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAV07752 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 20; Conservative
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV07752;
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Example 7; Page 30; Slpp; English

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                        containing compounds. It comprises contacting the phosphorus-containing compound which a 1,2,4-dithiazolidine-2,5-dione compound or a 3-substituted-1,2,4-dithiazolin-5-one compound. The method is especially useful for incorporation of phosphorothioate linkages into biologically important molecules such as DNA, RNA and phosphopeptides. Molecules containing such linkages are useful e.g. as antisense compounds for inhibiting gene expression, as reagents for studying DNA-protein or RNA-protein interactions, or as catalytic RNA. The present sequence represents an oligonucleotide with phosphorothioate linkages prepared by the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-HTLV-1 anti-sense oligo:nucleotide - is complementary to region of tax gene from human T-cell lymphotropic virus type 1 and inhibits viral antigen expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense; complementary; tax gene; inhibit; HTLV-1;
human T-cell lymphotropic virus type 1; viral antigen expression; ss.
                                                                                                                                                                                                                                                                                                                            Gaps
                present invention provides a method for sulphurising phosphorus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                   Length 20;
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                                                                                                                                                                                                                                             Sequence 20 BP; 1 A; 0 C; 0 G; 0 T; 19 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anti-HTLV antisense reference oligonucleotide HT.
                                                                                                                                                                                                                                                                                   Score 20; DB 1; L
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                  100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SOYA-) SOYAKU GIJUTSU KENKYUSHO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 8; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                  2708 TAAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                    20 TAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bb.
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                                                                                                                                                                                                                                                                                      0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT63649 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-AUG-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT63649;
                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 41
AAT63649/
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ð 셤 Oligonucleotide separated by capillary affinity gel electrophoresis.

(first entry)

04-JUN-1998

AAT86606;

gel electrophoresis; separation; polymer-gel;

Capillary afinity of polyacrylamide; ss.

W09745721-A1

Synthetic.

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Mycobacterium vaccae polypeptides - used to develop products for use in detection, therapy and prevention of mycobacteria infections or as immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This oligonucleotide is used in the DNA cloning strategies of the Mycobacterium vaccae antigens. The invention provides M. vaccae polypeptides that comprise an immunogenic portion of a soluble M. vaccae antigen, or a variant, where the antigen induces an immune response in patients previously exposed to a mycobacterium. Such M. vaccae patients previously exposed to a mycobacterium. Such M. vaccae polypeptides can be used in methods for enhancing non-specific immune response. The methods and products can be used for the detection, treatment and prevention of infectious diseases caused by mycobacteria such as M. vaccae, M. avium or M. tuberculosis. The products also have the ability to induce cell proliferation and cytckine products also have interferon-gamma and interleukin-12 production) in T cells, NK cells, interferon-gamma and interleukin-12 production) in T cells, B cells, Gr use in vaccines or immunotherapy of infectious diseases and cancers
                                                                                                                                                                                                                                                                                 Mycobacterium vaccae; antigen; therapy; prevention; cytokine production; M. avium; M. tuberculosis; immune response enhancer; cell proliferation; mycobacteria infection; vaccine; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prestidge RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 20; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tan P, Hiyama J, Visser E, Skinner MA, Scott LM,
                                                                                                                                                                                                                                               M. vaccae antigenic sequence hybridising oligo AD12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 99; 153pp; English.
           2709 AAAAAAAAAAAAAAAA 2728
                                              20 AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENE-) GENESIS RES & DEV CORP.
                                                                                                                                    AAV34591 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97WO-NZ000105.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-00705347.
                                                                                                                                                                                                          25-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.7'
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                         Mycobacterium vaccae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-216926/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          response enhancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-1996;
12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                            WO9808542-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                              05-MAR-1998
                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                        AAV34591;
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Separation of electrically charged target molecules - by capillary affinity gel electrophoresis using polymer-gel to which receptors for target molecules are bound.

Natt F;

Muscate A, Paulus A, (NOVS ) NOVARTIS AG.

WPI; 1998-041763/04.

97WO-EP002647.

23-MAY-1997; 24-MAY-1996;

04-DEC-1997.

96CH-00001320.

Example D3; Page 25; 41pp; English.

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Amixture of oligonucleotides (AAT86604-7) were separated by a new process using capillary affinity gel electrophoresis. The invention relates to selective separation of electrically charged target molecules in an analytical mixture. It comprises capillary affinity gel electrophoresis using a capillary tube which is at least partly filled with a polymer gel. Receptors for target molecules are covalantly bound to the polymer. An electric field of at least 50 volts/cm is applied. The capillary tube is charged with the analytical mixture. In a first separation stage, the target molecules in the mixture are bound to the receptors and the remaining components are eluted, optionally whilst coptionally in stages, so that the affinity of the target molecules are cluted and optionally in stages, so that the affinity of the target molecules for the receptor is eliminated and the target molecules are eluted and detected, optionally whilst splitting open. The process is useful for selective separation and/or determination of faraged organic compounds, such as oligonucleotides, peptides or carbohydrates. It may be used, e.g. for isolation of specific proteins and DNA molecules, purification of antibiotors. The process achieves higher resolution and selectivity than prior art processes, especially in the case of complex biological multitudes. The derivatised polymers may be synthesised specifically using standard methods
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0
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100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX27533 standard; RNA; 20 BP.
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AAX27533/c
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Gaps

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Pred. No. 6.8e+02; ; Mismatches 0; Indels

8 a AAT86606 standard; DNA; 20 BP.

AAT86606/c ID AAT8660 RESULT 418

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RESULT 421
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                internation retailed to allyloxymethyl halides (II). (I) are internation retailed to a solivy procected to a solivy the solivy procected to a solivy oxymethyl halides (II). (I) are intermediates for synthesis of RNA-oligonucleotides with predetermined nucleotide sequence, particularly by machine synthesis. The groups of specified above, apart from those on silyl, are those particularly for products in diagnosis, therapy, and as research tools, are well known, and are not dealt with in detail. (II) is an intermediate for (I). The silyloxymethyl halide reagent is easy to prepare, and yields are high. Incroduction of the silyloxymethyl group into the ribonucleoside is simple and rapid, and the acetal bond formed does not migrate, climinating particularly the prior art problem of 2' to 3' isomerisation. The methylenedioxy group spacer between the silyl group and nucleoside ring results in less steric hindrance than bulky direct silyloxy (I) in an analying first, a range of choices for the silyl substituents, to provide, e.g., acid or base stability; and second, higher yields in coupling Purer products are therefore obtained than in prior art, the coupling Purer products and longer chains of oligoribonucleotides to be a constituted of the coupling in the product of the coupling in the coupling is and longer chains of oligoribonucleotides to be considered to the coupling in the coupling in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                             2-Silyloxymethyl ribonucleosides and their phosphonate derivatives - have high purity, use in machine synthesis of ribonucleic acids, enable longer oligonucleotide chain construction, and larger amounts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention relates to silyloxymethyl protected D- or L-ribonucleosides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mycobacterium vaccae protein, antigen; T cell activation; cytokine; dendritic cell maturation; infectious disease; immune disorder; cancer;
                                                Silyloxymethyl; phosphonate; silyloxymethyl halide; diagnosis; ss; cyanoethyl phosphoramidate coupling; isomerisation; steric hindrance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                Synthetic RNA sequence produced by the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 20 BP; 0 A; 0 C; 0 G; 0 T; 20 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthesised successfully, and in shorter times
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mycobacterial 16S rRNA specific oligo AD12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 6; Page 25; 38pp; English.
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                                                                                                                                                                                                               98WO-EP005215.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.7
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                            Weiss PA,
                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-180963/15.
                                                                                                                                                                                                                                                                                    (PITS/) PITSCH S.
(WEIS/) WEISS P A.
(JENN/) JENNY L.
                                                                                                                                            WO9909044-A1.
                                                                                                                                                                                                                 17-AUG-1998;
                                                                                                                                                                                                                                                    L8-AUG-1997;
                                                                                                                                                                                25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                              Pitsch S,
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   8
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Pharmaceutical compositions comprising the proteins or nucleic acid beareaceutical compositions can be used for the treatment, prevention, and detection of disorders including infectious diseases, immune disorders and cancer. In particular, the compounds and methods are used for treatment of diseases of the respiratory system, such as mycobacterial infections, asthma, allergies, tuberculosis, leprosy, sarcoidosis and lung cancers, and disorders of the skin such as psoriasis, atopic dermatitis, eczema, allergic contact dermatitis, allopecia areata, and skin cancers such as basal carcinoma, squamous cell carcinoma and melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention provides heat-killed Mycobacterium vaccae, or recombinant M. vaccae proteins may be employed to activate T cells and natural killer cells, to stimulate the production of cytokines, to enhance the expression of co-stimulatory molecules on dendritic cells and monocytes, and to enhance dendritic cell maturation and function. The proteins can be expressed by standard recombinant methodology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
respiratory system; mycobacterial infection; allergy; tuberculosis; leprosy; sarcoidosis; lung cancer; asthma; skin disorder; psoriasis; dermatitis; eczema; alopecia areata; skin cancer; basal carcinoma; squamous cell carcinoma; melanoma; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Electrochemical detection; glucose; cholesterol; urea nitrogen; bilirubin; uric acid; haemoglobin; lactic acid; body fluid; blood; plasma; serum; urine; lymph diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prestidge RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tan P, Watson J, Visser ES, Skinner MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enhancing immune response to an antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Page 177; 243pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENE-) GENESIS RES & DEV CORP LTD
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97US-00997362.
98US-00095855.
98US-00156181.
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Matches 20; Conservative
                                                                                                                                                                                                   Mycobacterium vaccae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-430163/36.
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                                                                                                                                                                                                                                                                                                                                                                                      23-DEC-1998;
                                                                                                                                                                                                                                                                                                                            01-JUL-1999,
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                                                                                                                                                                      Synthetic
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This invention describes a novel method for quantitatively analysing a biochemical compound (I) which comprises contacting (I) with double atranded DNA fixed to the surface of an electrode at their terminals in which electrochemically active threading intercalators are intercalated, in an aqueous medium under application of electric potential to the clectrode in the presence of an oxidase which oxidizes the biochemical compound and becomes reduced, and detecting electric current flowing between the electrode and second electrode in the aqueous medium. The method is useful for detection of biochemical compounds such as plucose, cholesterol, urea nitrogen, bilirubin, uric acid, heameglobin and lactic acid in body fluids such as whole blood, plasma, serum, urine, and lymph for diagnosis of various diseases. The method allows detection of biochemical compounds quickly and easily with a high sensitivity using a simple apparatus. This sequence represents DNA fragment used as fixed probe DNA in the method of the invention
                                                                                                                                                                                                 Quantitative analysis of a biochemical compound such as glucose, in body a body fluid such as blood, comprising detecting enhanced electron transfer between an oxidase and a DNA-immobilized electrode, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stem cell factor; SCF; haematopoietic progenitor cell; blood forming; primitive progenitor cell; haematopoietic disorder; syngensic; allogeneic; autologous bone marrow marnsplant; gene therapy; transfection; haematopoietic stem cell; acute blood loss; neoplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stem cell factor universal oligonucleotide 220-7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                Example 1; Page 7; 14pp; English
                                                                                                                 Takagi M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 AAAAAAAAAAAAAAAA 1
                                                                    (FUJF ) FUJI PHOTO FILM CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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90US-00537198.
90US-00573616.
90WO-US005548.
90US-00589701.
      99JP-00001111.
99JP-00143599.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.(
Matches 20; Conservative
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                                                                                                               Ogawa M, Takenaka S,
                                                                                                                                                                                                                                                                   diagnosis of disease.
                                                                                                                                                          WPI; 2000-444372/39.
         06-JAN-1999;
                           24-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-OCT-1989;
11-JUN-1990;
24-AUG-1990;
28-SEP-1990;
01-OCT-1990;
04-OCT-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA13753;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 423
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      셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for quantitatively analysing a biochemical compound (1) which comprises contacting (1) with double stranded DNA fixed to the surface of an electrode at their terminals in which electrochemically active threading intercalators are intercalated, in an aqueous medium under application of electric potential to the electrode in the presence of an oxidase which oxidizes the biochemical compound and becomes reduced, and detecting electric current flowing between the electrode and a second electrode in the aqueous medium. The method is useful for detection of biochemical compounds such as glucose, cholesterol, urea nitrogen, bilirubin, uric acid, haemoglobin and lactic acid in body fluids such as whole blood, plasma, serum, urine, and lymph for diagnosis of various diseases. The method allows detection of biochemical compounds quickly and easily with a high sensitivity using a biochemical compounds.
                                                                                                                                                                                                                                                                                                                                                                       Quantitative analysis of a biochemical compound such as glucose, in body a body fluid such as blood, comprising detecting enhanced electron transfer between an oxidase and a DNA-immobilized electrode, useful for diagnosis of disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . This sequence represents DNA fragment used as a target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Electrochemical detection; glucose; cholesterol; urea nitrogen;
bilirubin; uric acid; haemoglobin; lactic acid; body fluid; blood;
plasma; serum; urine; lymph diagnosis; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Electochemical detection method fixed probe DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sample in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AAAAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                   Takenaka S, Takagi M;
                                                                                                                                                                                                                                     (FUJF ) FUJI PHOTO FILM CO LTD.
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                                                                                                                             07-JAN-2000; 2000EP-00100126.
                                                                                                                                                                          99JP-00001111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.
                                                                                                                                                                                                                                                                                                                              WPI; 2000-444372/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  simple apparatus
                                           EP1018646-A2
                                                                                                                                                                        06-JAN-1999;
                                                                                                                                                                                               24-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP1018646-A2
                                                                                    12-JUL-2000
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Synthetic.
                                                                                                                                                                                                                                                                                     Ogawa M,
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ID AAA404
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Matches

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Gaps .. 0

0; Indels

Tue Nov

Production of hematopoietic cells suitable for administration to a subject using progenitor cells and expanding the cells using stem cell WPI; 2000-259135/23 (AMGE-) AMGEN INC. Zsebo KM, 

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Amethod has been developed of making haematopoietic cells suitable for administration to a subject. The method comprises: (a) obtaining cadministration to a subject. The method comprises: (a) obtaining to administration to a subject. The method comprises: (a) obtaining to be adding to the cells a haematopoietically effective dose of a polypeptide product having at least part of the primary structural confirmation and one or more of the biological properties of naturally occurring stem cell factor (SCF). The method is useful for stimulating primitive progenitor cells including early haematopoietic progenitor cells including early haematopoietic progenitor cells including early haematopoietic cells of both myeloid and lymphoid lineages. Increases in haematopoietic cells of both myeloid and lymphoid lineages. Cor expanding early haematopoietic progenitors in syngeneic, allogeneic cells of cor autologous bone marrow transplant. SCF is useful for enhancing the cells. Cor autologous bone marrow transplant. SCF is useful for enhancing the myelosuppressive effects of anti-HIV drugs such as AZT and for enhancing haematopoietic recovery cafter acute blood loss and as a boost to the immune system for fighting continuous and continuous continuous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide which is used in an example from the present invention
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Example 3; Fig 12C; 123pp; English.
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Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels Similarity 100.0%; Score 20; LL. Similarity 100.0%; Pred. No. 6.8 CTAAAAAAAAAAAAAA 2726 CTAAAAAAAAAAAAAAA 1 Local Sim 2707 20 Query Match Matches ò g

Oligonucleotide #5 for conjugation to abietane derivative. AAZ91117 standard; DNA; 20 (first entry) 06-JUN-2000 AA291117; RESULT 424 

Abietane derivative; labelling; diagnostic test; biotin substitute; ss.

FR2781802-A1 Synthetic.

04-FEB-2000

98FR-00010084 31-JUL-1998;

98FR-00010084 31-JUL-1998;

(INMR ) BIO MERIEUX.

Delair T, Mandrand B; Veron L, Battail PN, Charles MH, Piga N, WPI; 2000-239603/21

Saturated and unsaturated derivatives of abietic acid and their conjugated derivatives with natural and synthetic polymers, having use in diagnostics, chemical reactions and analysis.

Example 5; Page 20; 39pp; French

Martin FH;

Bosselmann RA,

Suggs SV,

derivatives. The new compounds may be used directly or indirectly in the development of new diagnostic tests, to follow infections, especially viral infections, to follow and/or measure chemical products, especially potential pollutants. In diagnostic tests they may be used as markers, or to form a universal solid phase after immobilization on a solid support, to produce monoclonal antibodies or polyclonal antibodies having diagnostic uses. The oligomuclectides AAZ91113-Z91117 represent examples of sequences that can be labeled with the new abietane derivatives. The but because they are not found naturally in humans the risk of potential interactions with biological molecules is eliminated The invention relates to novel saturated and unsaturated abietane 

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; rative 0; Mismatches 0; Indels 20; Conservative Local Similarity Query Match Matches

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AAA50193 standard; DNA; 20 07-NOV-2000 (first entry) AAA50193; 

Phosphodiester oligonucleotide; H-phosphonate chemistry; 88. 2'-Methoxyethoxy-modified oligonucleotide.

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Gaps

.; 0

/notes "2'-methoxyethoxy modified thymidine" Location/Qualifiers Ø /\*tag= Key modified\_base Synthetic.

11-FEB-2000; 2000WO-US003543. WO200047593-A1 17-AUG-2000

Maier MA; (ISIS-) ISIS PHARM INC. Manoharan M,

99US-00250075

12-FEB-1999;

WPI; 2000-558188/51.

Preparation of mixed backbone oligomeric compounds useful as e.g. primers for diagnostic tests, involves oxidation of H-phosphonate internucleoside linkages to phosphodiester internucleoside linkages.

Example 12; Page 34; 49pp; English.

The present sequence is that of a phosphodiester oligonucleotide containing 20 T nucleobases, 19 having a 2'-methoxyethoxy group on its 5' ribosyl sugar moiety. It is an example of an oligomeric compound produced according to the methods of the invention. The invention provides

7 10:41:34 2006

Tue Nov

it and thereby act as functional modifiers of immunostimulatory

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compounds and methods for the preparation of mixed backbone oligomeric, addition to phosphoratioate and/or phosphoramidate internucleoside linkages in linkages. The methods also include incorporation of boranophosphate internucleoside linkages. The methods utilise H-phosphonate intermediates internucleoside linkages. The methods utilise H-phosphonate intermediates phosphonate internucleoside linkages. Each contiguous regions of 1 or more H-phosphonate internucleoside linkages. Each contiguous region is subsequently oxidized to phosphodiester, phosphorothicate, phosphoromidate or boranophosphate internucleoside linkages prior to this phosphoromidate or boranophosphate internucleoside linkages prior to this phosphoremic are prepared in the internucleoside linkages prior to this phosphoremic phosphoremic phosphoremic phosphate internucleoside linkages prior to this phosphoremic p
                                                                                                                                                                                                                                                                         this manner by oxidizing adjacent regions with different reagents.

Oligomeric compounds of the invention are prepared using novel oxidation steps that oxidize a region of 1 or more H-phosphonate internucleoside linkages without degrading existing linkages that have been previously pxobes, The oligonucleotides obtained are useful as primers in PCR, probes, linkers, gene fragments and for other diagnostic tests on e.g. biological tissue, fluid, cells etc., as research reagents, and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory oligodeoxynucleotide; immunostimulatory ODN; immunostimulatory DNA-binding protein; nucleolin; hnRNP D; AUF1; hnRNP AI, lupus La protein; functional modifier identification; agonist; antagonist; mim.c; inhibitor; drug screening; cellular target identification; oligonucleotide optimisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Phosphorothioate poly T oligonucleotide, SEQ ID NO:17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Scc...
100.0%; Prev
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03-MAR-2000; 2000US-0186845P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               antiviral agents
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09-MAR-2001
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Oligodeoxynucleotide (ODM) activity. Such modifiers of ODM activity consist of immunostimulatory DNA binding inhibitors, immunostimulatory DNA binding inhibitors, immunostimulatory DNA agonists and antagonists.

Immunostimulatory DNA-binding proteins can also be used in screening methods to identify immunostimulatory DNA binding competitors, and to optimize an immunostimulatory ODM for immune stimulation. Isolated complexes of an immunostimulatory DNA-binding protein bound to an immunostimulatory ODN can additionally be used to screen a panel of candidate target molecules to identify the cellular target molecules of the immunostimulatory DNN-binding proteins used in the methods of the invention are the RNM-binding proteins cublewen immunostimulatory DNA and an immunostimulatory DNA-binding proteins methods are useful for identifying a compound that inhibits interaction between immunostimulatory DNA and an immunostimulatory DNA-binding complex is useful in screening consists useful in immunotherapy. The complex is useful in screening to identifying agonists useful in immunotherapy. The complex is useful in screening consisting immunostimulatory DNA cellular target molecules. The candidate immunostimulatory DNA cellular target consisting immunostimulatory DNA cellular target consisting immunostimulatory consisting and for identifying activity relationships of immunostimulatory consisting immunostimulatory consisting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA-binding proteins and immunostimulatory ODNs. The present sequence represents an oligonucleotide used in an exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. ...
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(IOWA ) UNIV IOWA RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 AAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC87230 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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$$$$$$$$$$$$$$$$$$$$$$$$$$$
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The invention relates to the use of an immunostimulatory single-stranded DNA-binding protein in screening assays to identify compounds which bind

Example 1; Page 45; 95pp; English.

Immunostimulatory DNA binding proteins to identify immunostimulatory functional modifiers, immunostimulatory DNA binding competitors and toptimize immunostimulatory oligodeoxynucleotides for stimulation.

WPI; 2001-016002/02.

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is useful in screening for immunostimulatory DNA cellular target molecules. The candidate immunostimulatory ODN competitors allow the investigation of structure/activity relationships of immunostimulatory DNA-binding proteins and immunostimulatory ODNs. The present sequence represents an oligonucleotide used in an exemplification of the invention Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; Ative 0; Mismatches 0; Indels 2709 AAAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAAA 1 Query Match Best Local Similarity 100.C Warcheg 20; Conservative ð 요

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Gaps ö

> AAC87241 Standard; DNA; 20 BP 09-MAR-2001 (first entry) AAC87241; RESULT 428 A CONTRACTOR OF A CONTRACTOR O

Poly T oligonucleotide, SEQ ID NO:20.

Immunostimulatory oligodeoxynuclectide; immunostimulatory ODN; immunostimulatory DNA-binding protein; nucleolin; hnRNP D; AUF1; hnRNP Ah; lupue La protein; functional modifier identification; agonist; antagonist; mimic; inhibitor; drug screening; ellular target identification; oligonucleotide optimisation; immunotherapy; 88.

Synthetic.

WO200067023-A1.

28-APR-2000; 2000WO-US011697. 09-NOV-2000

29-APR-1999; 99US-0131830P. 03-MAR-2000; 2000US-0186845P.

CPGI-) CPG IMMUNOPHARMACEUTICALS GMBH. (IOWA ) UNIV IOWA RES FOUND.

Noll BO,

Krieg AM;

WPI; 2001-016002/02.

Immunostimulatory DNA binding proteins to identify immunostimulatory  ${\tt DN}$  functional modifiers, immunostimulatory  ${\tt DNA}$  binding competitors and to

The invention relates to the use of an immunostimulatory single-stranded CC DNA-binding protein in screening assays to identify compounds which bind DNA-binding protein in screening assays to identify compounds which bind CC to it and thereby act as functional modifiers of immunostimulatory CC consist of immunostimulatory DNA binding inhibitors, immunostimulatory DNA agonists and antagonists.

CC manunostimulatory DNA-binding proteins can also be used in screening centends to identify immunostimulatory DNA binding proteins can also be used in screening complexes of an immunostimulatory ODN for immune stimulators, and to optimize an immunostimulatory ODN can additionally be used to screen a panel of immunostimulatory ODN can additionally be used to screen a panel of candidate target molecules to identify the cellular target molecules of the immunostimulatory ODN. The immunostimulatory DNA-binding proteins cused in the methods of the invention are the RNA-binding proteins cused in the methods of the invention are the RNA-binding proteins contoled in, hnRNP D, AUPI, hnRNP AI and lupus La protein. The screening methods are useful for identifying a compound that inhibits interaction between immunostimulatory DNA and an immunostimulatory DNA-binding complexed the protein and for identifying a compound that inhibits interaction between immunostimulatory DNA and an immunostimulatory DNA-binding complexed the information and information in immunotherapy. The complex complexed the information in immunotherapy in the complexed that interaction in the method in the immunostimulatory DNA and an immunostimulatory DNA-binding complexed the immunostimulatory DNA and an immunostimulatory DNA-binding complexed the immunostimulatory DNA and an immunostimulatory DNA-binding complexed the immunostimulatory DNA-bindi ö represents an oligonuclectide used in an exemplification of the invention processing and the conditions in mannostimulatory DNA cellular target is useful in screening for immunostimulatory DDN competitors allow the molecules. The candidate immunostimulatory ODN competitors allow the investigation of structure/activity relationships of immunostimulatory NNA binding nyreleins and immunostimulatory ODNs. The present sequence Gaps optimize immunostimulatory oligodeoxynucleotides for stimulation. ö / Match 0.7%; Score 20; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 6.8e+02; nes 20; Conservative 0; Mismatches 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; DNA-binding proteins and immunostimulatory ODNs. 2709 AAAAAAAAAAAAAAAAA 2728 Example 1; Page 45; 95pp; English Query Match Matches ð g

20 AAAAAAAAAAAAAAAAAA 1

BP.

AAS10402 standard; DNA; 20

RESULT 429 AAS10402, AAS10402;

3' RNA end labeling; DNA template; Okazaki fragment; 5' overhang; ss. DNA template for 3' end labeling of an RNA molecule, #14. 98US-00173936. 97US-0063757P 24-OCT-2001 (first entry) Huang Z, Szostak JW; (CHEN/) CHEN G. (HUAN/) HUANG Z. (SZOS/) SZOSTAK J W. 16-OCT-1998; US6238865-B1. 17-OCT-1997; 29-MAY-2001. Synthetic. 

Modifying a 3' terminus of a pre-selected DNA sequence, useful for labeling and modifying 3'-termini of other nucleic acids, comprises using a synthetic nucleotide template with a defined overhang nucleotide. WPI; 2001-366470/38.

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Example 5; Col 13; 22pp; English
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The sequence represents a synthetic DNA template molecule used to demonstrate the method of the invention. The invention relates to a method of modifying (e.g. 3' end labelling with 32P dATP) the 3' terminus of an RNA molecule by providing a DNA oligonucleotide, complimentary to the 3' end of the RNA molecule, with an overhang at the 5' end which allows incorporation of the labelling nucleotide into the RNA molecule. The method, based on the synthesis of Okazaki fragments, is useful for Tabeling and modifying the 3'-termini of other nucleic acids such as DNA fragments. The method is a simple and efficient way of labelling or modifying RNA 3'-termini using DNA polymerase and a synthetic template with defined overhang nucleotides

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

Gaps ; 0 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; /ative 0; Mismatches 0; Indels 2709 AAAAAAAAAAAAAAAA 2728 20; Conservative Query Match Best Local Similarity Matches ð g

AAD16997 standard; DNA; 20 BP 29-NOV-2001 (first entry) Capture probe CP5' AAD16997; RESULT 430 AAD16997 

Scaffold protein; antibody mimic; fibronectin type III domain; randomised loop; randomised beta-sheet; diagnostic purpose; protein designing; probe; tenth module of human Fn3; 10Fn3; fibronectin module of type III; Fn3; ss.

Unidentified

WO200164942-A1. 07-SEP-2001 28-FEB-2001; 2001WO-US006414

29-FEB-2000; 2000US-00515260

(PHYL-) PHYLOS INC.

Kuimelis RG; Wagner RW, Lipovsek D,

WPI; 2001-557782/62.

Fibronectin scaffold protein array for obtaining a protein/compound which binds to a compound/protein, comprises a fibronectin type III domain having a randomized loop, a randomized beta-sheet or their combination. Disclosure; Page 41; 67pp; English.

The present invention relates to an array of proteins (antibody mimics) comprising a fibronectin type III domain having a randomised loop, a randomised beta-sheet, or their combination, and has the capacity to bind to a compound that is not bound by a corresponding naturally-occurring tibronectin, immobilised onto a solid support. The antibody mimics is useful for detecring a compound preferably a protein, in a biological sample. It is also useful to detect one or more different analytes simultaneously in a sample. Hence is useful for diagnostic purposes. It is also useful for the purpose of designing proteins capable of binding to virtually any compound of interest. The present sequence is a capture probe used to self-assemble and anchor the tenth module of human

Gaps fibronectin module of type III (Fn3) (10Fn3) which is used in an exemplification of the invention ö Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; 100.0%; Pred. No. 2709 AAAAAAAAAAAAAAAA 2728 0.7%; 20; Conservative Similarity 20 Query Match Best Local Matches

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BP. AAF60896 standard; DNA; 20 (first entry) 15-MAY-2001 AAF60896; AAF60896

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RESULT 431

Conjugate forming oligonucleotide ONS SEQ ID 5.

Transport; membrane; cytostatic; virucide; vasotropic; dermatological; antipsoriatic; antiasthmatic; gene therapy; tumor cell; antisense; tumor therapy; drug; phosphodiester linkage; ss.

Unidentified

DE19935302-A1.

08-FEB-2001.

99DE-01035302 28-JUL-1999; 99DE-01035302. 28-JUL-1999; (AVET ) AVENTIS PHARMA DEUT GMBH

Schwerdel M;

Gothe G,

Unger E,

Greiner B,

Jhlmann E,

WPI; 2001-203679/21

New substituted aryl conjugates of parent molecules, especially oligonuclectides, having improved transmembrane and intracellular transport properties, useful as medicaments or diagnostic agents.

Disclosure; Page 9; 28pp; German.

This invention describes a novel conjugate (I) which consists of (A) a molecule to be transported and (B) at least one aryl residue of formula - CC molecule to be transported and (B) at least one aryl residue of formula - CC Ar-(K-CY)-R I) n (II). Ar = group containing at least one aromatic ring; CX = O or N (Bic); Y = O, SO NH-R\_2 (Bic); R I = optionally substituted containing double and/or triple bonds); R\_2 = optionally substituted 1.18C alkyl (optionally containing double and/or criple bonds); n = integer of 1 or more. (A) is bonded to (B) directly or criple bonds); n = integer of 1 or more. (A) is bonded to (B) directly or criple bonds); n = integer of 1 or more. (A) is bonded to (B) directly or criple bonds; n = integer of 1 or more. (A) is bonded to (B) at least one aryl residue (not restricted to compercial so (i) the preparation of a conjugate (I) of (A) a molecule to be transported and (B) at least one aryl residue (not restricted to compercial (B) is to be bonded, preparing (B) and reacting (A) and (B); at which (B) is to be bonded, preparing (B) and reacting (A) and (B); cand (ii) the use of aryl groups (II) (optionally bonded via a chemical group) for transporting (A) across biological membranes. The products of the invention have cytostatic, virucide, vasotropic, dermatological, carriage functionally bacterial, yeast or mammalian cells, including human cells, cyparticularly tumor cells). Medicaments, diagnostic agents and test kits containing (I) are also claimed. Typically (I) are antisense 

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cell interactions (e.g. restencess, vitiligo, psoriasis or asthma); or labeled oligonucleotides for in vivo diagnostic use, e.g. by in situ bybridization. Conjugation with (B) markedly improves the cellular uptake of (A), e.g. in tumor cells. (B) include fluorescein derivative residues, in which case the conjugates (I) are fluorescently labeled, allowing microscopic monitoring of cellular uptake etc. The cellular uptake of (I) is superior to that obtained using other conjugated groups related (II); e.g. oligonucleotides conjugated with fluorescein diacetate (within the scope of (B)) have superior uptake to corresponding fluorescein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide-nanoparticle probe, diagnostic, forensic analysis, nucleic acid detection, nanostructure, biochip, biofilter, drug delivery,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         having at least 2 portions, comprising treatment with nanoparticles that carry oligonucleotides complementary to at least 2 parts of [1], where detectable change caused by hybridisation of the oligonucleotide to [1] is observed. The method is used to detect (or to separate) specific [1], e.g. for diagnosing a wide variety of diseases, sequencing, in forensic
treating viral infections or diseases associated with integrins or cell-
                                                                                                                                                                                                                                                                         Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the invention relates to a method for detection of nucleic acid (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting and separating nucleic acid, useful e.g. for diagnosis, comprises reaction with nanoparticles that carry oligonucleotides complementary to parts of the target.
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                                                                                                                                                                                                                                      DB 1; Length 20; 6.8e+02;
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                                                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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0
                                                                                                                                                                                                                                    0.7%; Score 20; DB
100.0%; Pred. No. 6.8
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide-nanoparticle probe #52.
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                                                                                                                                                                                                                                                                                                                            AAS63428 standard; DNA; 20 BP
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2000US-00603830.
2000US-0213906P.
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Park S, Li Z;
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                                                                                                                                                                                                                                                                         20; Conservative
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                                                                                                                                                                                                                                                         Local Similarity
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26-JUN-2000;
26-JUN-2000;
08-DEC-2000;
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                                                                                                                                                                          conjugates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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The invention relates to a method for enrichment of natural antisense messenger RNA. This method involves generating a population of CDNA from mRNA, incubating the generated CDNA to produce double stranded hybrid DNA molecules consisting of sense and antisense CDNA, treating the hybrid colecules using DNA polymerase with an exonuclease activity, amplifying the double stranded molecule using a deoxythymidine (dT) primer and cloning the amplified double stranded molecule. This method is useful for enrichment of natural antisense mRNA from any natural source of RNA. It is used to detect whether mRNAs have a natural anti-sense counterpart. The method provides a basis for finding new genes with important cellular cyclular provides a transfal for development of an antisense-based provides a starting material for development of an antisense-based the sense gene or transcript is involved. The present sequence is KU\_REV analysis etc., and generally to detect analytes other than (I). The oligonucleotide-derivatised nanoparticles are also useful for preparing nanostructures useful, for example, as biochips, biofilters, mechanical devices, separation membranes, chemical sensors, in computers, and for drug delivery. Very stable nanoparticle-oligonucleotide conjugates can be produced, allowing their direct use (as probes) in polymerase chain reaction, i.e. they survive multiple heating/cooling cycles so do not need to be added after amplification. (I) are detected by simple colour change, without the need for special equipment, making possible rapid field testing for e.g. pathogens. Assi314-AAS65448 represent oligonucleotide-nanoparticle probes, and related sequences, used in the Enrichment and detection of natural antisense mRNA comprises generating double stranded hybrid cDNA using a polymerase with an exonuclease activity, amplifying using a DT primer and cloning. Gaps natural antisense mRNA enrichment; antisense-based therapy; .. 0 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; :ive 0; Mismatches 0; Indels Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; Human ku autoantigen amplifying KU\_REV primer. 2709 AAAAAAAAAAAAAAAA 2728 primer; ku autoantigen; ss. 1 AAAAAAAAAAAAAAA 20 Example; Page 12; 37pp; English. Grossman A; BP. 06-OCT-2000; 2000WO-US027557. 99US-0157843P AAD03632 standard; DNA; 20 19-JUN-2001 (first entry) (QUAR-) QUARK BIOTECH INC Best Local Similarity 100. Matches 20; Conservative method of the invention WPI; 2001-266326/27 Einat P, WO200125488-A2 06-OCT-1999; Homo sapiens. 12-APR-2001. AAD03632; Gilad S, Query Match Human; RT-PCR 433 AAD03632 RESULT 222222222222222 ò g ö

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a nucleic acid having at least 2 portions. The method comprises thybridising the nucleic acid with oligonuclectides, such as the present sequence, attached to a substrate and/or particle and detecting a change in colour, conductivity or optical density. The method is useful for the diagnosis and/or monitoring of diseases, in forensics, in DNA sequencing, for paternity testing, for cell line authentication and for monitoring gene therapy. Detecting nucleic acids based upon observing a colour change is cheap, fast, simple, and does not require specialised or expensive equipment. The nanoparticle oligonucleotide conjugates remain stable for at least 6 months. A single base mismatch and as little as 20 femtomoles (fM) of target can be detected using the conjugates
reverse transcription PCR (RT-PCR) primer used for amplifying human ku autoantigen sequence. This primer is used in endogenous antisense identification (EASI) procedure for enrichment of natural antisense mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is an oligonucleotide used in a method for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid detection; nanoparticle-oligonucleotide conjugate;
disease diagnosis; forensic analysis; DNA sequencing; paternity testing;
cell line authentication; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detecting nucleic acid, useful for e.g. diagnosis of diseases, forensics and DNA sequencing, comprises observing detectable change brought about by hybridization of nucleic acid with substrate or particle bound oligonucleotides.
                                                                                                                                        Gaps
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                                                                                                       Length 20;
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                                                                    Sequence 20 BP; 6 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                   Score 20; DB :
Pred. No. 6.86
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                          Random oligonucleotide, SEQ ID NO: 53
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                                                                                                                                                                      2010 TCATGCCAACTCCAGAGCAG 2029
                                                                                         0.7%; Scc_
100.0%; Prev
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                                                                                                                                                                                      1 TCATGGCAACTCCAGAGCAG
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26-APR-2000; 2000US-0200161P.
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                                                                                                                   Best Local Similarity 100.
Matches 20; Conservative
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(LETS/) LETSINGER R L.
(MUCI/) MUCIC R C.
(STOR/) STORHOFF J J.
(ELGH/) ELGHANIAN R.
(TATO/) TATON T A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                                                                     Query Match
                                                                                                                                                                                                                                                        RESULT 434
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ស្ត្រប្តប្តវត្ត
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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The sequence represents a cyclic disulphide linked oligonucleotide which may be coupled with colloidal gold particles (nanoparticles) and used to demonstrate the method of the invention. The invention relates to isolating or detecting a nucleic acid of interest, in a mixture of nucleic acids, by binding it to 2 or more complimentary nucleotides which nucleic acids, by binding it to 2 or more complimentary nucleotides which have a nanoparticle attached to their 5' ends. The nanoparticles (e.g. colloidal gold) are used to both isolate and detect (e.g. by linking the particle to a fluorescent probe) the resultant complex. The methods are useful for detecting nucleic acids, natural or synthetic, and modified or unmodified. The methods may also be applied in the disagnosis of genetic, bacterial and viral diseases, in forenaics, in DNA sequencing, for cell line authentication, and for monitoring gene therapy. The methods are further useful in research and analytical laboratories in DNA sequencing, in the field to detect the presence of specific pathogens, for quick identification of an infection to assist in drug prescription, and in homes and health centres for inexpensive first-
                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting a nucleic acid useful in e.g. diagnosing genetic, bacterial or viral diseases, by contacting the nucleic acid with oligonucleotides attached to nanoparticles and having sequences complementary a portion of the nucleic acid.
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/note= "A is covalently linked to a cyclic-disulphide
moiety"
                                              Gaps
                                                                                                                                                                                                                                                                                                                                      Nanoparticle, cyclic disulphide-oligonucleotide, DNA detection, DNA isolation; genetic disease; bacterial disease; viral disease; forensic science; paternity testing; gene therapy; ss.
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            Length 20;
                                            0; Indels
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          DB 1; Le
0.7%; Scor.
100.0%; Pred. No. c.
0; Mismatches
                                                                                                                                                                                                                                                                                                      Oligonucleotide-cyclic disulphide linker, d.
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                                                                              2709 AAAAAAAAAAAAAAAAA
                                                                                                                1 AAAAAAAAAAAAAAAAA
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26-APR-2000; 2000US-02001G1P.
26-JUN-2000; 2000US-0060830.
12-JAN-2001; 2001US-00766590.
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                                                                                                                                                                                                  AAS10371 standard; DNA; 20
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                                            Conservative
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                          Similarity
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misc_feature
          Query Match
Best Local Simi
Matches 20;
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                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                    AAS10371;
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                                                                                                                                                                  RESULT 435
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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Thi immune response and to activate immune cells. Note: the
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line screening. The methods, which are based on observing colour change with the naked eye, are cheap, fast, simple, robust (reagents are stable), do not require specialised or expensive equipment, and little or no instrumentation is required
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccinating against tumors, infectious diseases, allergies and asthma
using immunostimulatory Py-rich and TG nucleic acids.
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                                                                                                                                          Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
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                                                                                                     Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                          0.7%; Score 20; DB
100.0%; Pred. No. 6.8
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                 AAF99427 standard; DNA; 20 BP
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                                                                                                                                              Query Match
Best Local Similarity 100.(
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-273485/28
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DB 1; Length 20;

0.7%; Score 20;

Query Match

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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immunose. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, chaemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is slowered also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a that to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothicate backbone
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using immunostimulatory Py-rich and TG nucleic acids.
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100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                            Indels
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           Pred. No. 6.8e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                    Immunostimulatory nucleic acid #215.
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                                                                2709 AAAAAAAAAAAAAAAA 2728
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100.08; Pre-
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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(COLE-) COLEY PHARM GMBH.
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Best Local Similarity 100.
Matches 20; Conservative
             Best Local Similarity 100.
Matches 20; Conservative
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Universal stem cell factor (SCF) related oligonucleotide SEQ ID NO:33.
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                                                                                                                                                                                                                                                                                                                                                                  into cell in vitro.
                                                                                                                                                                                                                                                             (AMGE-) AMGEN INC.
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                                                                                   US6207454-B1
                                                                                                                                                                                                    25-NOV-1992;
21-DEC-1993;
24-MAY-1995;
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24-AUG-1990
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                                                             Synthetic.
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Matches
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                                                                                                                 Vaccine, cytostatic, virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory, tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid can be pyrimidated (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia acid and/or staphylococus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a first to a Thi immune received to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                 Vaccinating: against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present sequence may have a phosphorothicate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Prea....
                                                                                            Immunostimulatory nucleic acid #547.
                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 101; Þage 49; 338pp; English.
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                         BP.
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27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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                        AAF99431 standard; DNA; 20
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                                                                     (first entry)
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les 20; Conserv
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                                              AAF99431;
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RESULT 438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mammalian cell in vitro, involves exposing cell that expresses a stem
cell factor receptor to stem cell factor, and introducing polynucleotide
Stem cell factor, SCF; stem cell factor receptor; blood cell disorder; gene therapy; PCR primer; mutagenesis; probe; ss.
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100.0%; Pred. No. 6.8e+02;
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90US-00537198.
90US-00573616.
92US-0088225.
93US-00172329.
95US-0049653.
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US6169176-B1.
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02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
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                  Blumberg B;
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                                                                                                                                                                                                                                                                        The present invention relates to a method for preparing phosphorothioate oligonucleotides having at least one nucleoside with a 2' modification. The method comprises phosphitylating the 5'-hydroxyl of a nucleic acid group having at least one nucleoside with a 2' modification in an acetonitrile. The present sequence was used to illustrate the method of the present invention. The method is useful for synthesising sulphurised 2' substituted phosphorothioate oligonucleotides, which may be used in molecular biological research, in applications such as anti-viral therapy, and for determining the stereochemical pathways of certain enzymas which recognise nucleic acids
                                                                                                                                                                                                                 Preparing sulfurized 2' substituted phosphorothioate oligonucleotides useful in biological research, comprises phosphitylating the 5'-hydroxyl of a nucleic acid having a nucleoside with a 2' modification.
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                  /mod_base= OTHER
/note= "All bases are phosphorothioate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid identification; DNA library screening; ss
                                              /*tag= b
/mod base= OTHER
/note= "Modified with 2'-O-methyl"
                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. ...
                                                                                                                                                                               Cheruvallath ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleotide sequence of a cDNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                        Example 23; Col 11; 7pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ААААААААААААААА 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-00454704
                                                                                                                        11-JAN-2000; 2000US-00481486
                                                                                                                                          97US-00950779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH78547 standard; cDNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                               Ravikumar VT,
                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                  WPI; 2001-407218/43
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 20; Conserv
 modified_base
                                       modified_base
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                                                                                    JS6242591-B1
                                                                                                                                           15-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2001
                                                                                                      05-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparing novel deoxynucleic alkyl thiourea oligonucleotide for use in antisense therapy, by synthesizing oligonucleotides comprising backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Deoxynucleic S-Methythiourea; DNmt; antisense therapy; cardiovascular disease; inflammatory disease; neurocellular disease; antiviral therapy; human immunodeficiency virus; human-cytomegalovirus; influenza; herpes; infection; 88.
                                                                                                                                                                                               Screening nucleic acids (NA) in pool of interest comprises pooling, expressing NA to form expression product pool and identifying NA in NA pool corresponding to expression product pool having interaction with target moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 22; 19pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0091481P.
98US-0111800P.
99US-00347443.
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Best Local Similarity 100.8
Matches 20, Conservative
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(REGC ) UNIV CALIFORNIA
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                                                                                                                                       WPI; 2001-588900/66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bruice
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The present sequence was used to demonstrate the ability of deoxynucleic S-Methythiourea (DNmt) compounds to form triplexes with DNA oligomers. An increase in the C content of the oligos resulted in a large decrease in the C content of the oligos resulted in a large decrease in binding. This experiment was performed as an example of a method for preparing oligonucleotides comprising a backbone of alkyl or alkoxy comprising on a performed in second to a large therapy, to inhibit production of proteins associated with genetic diseases, cardiovascular, inflammatory and neurocellular diseases, and for antiviral therapy, e.g. to treat comman immunodeficiency virus, human-cytomegalovirus, influenza and herpes infections. The compounds are also useful as diagnostic reagents to detect the presence or absence of the target DNA or RNA sequences to detect the presence or absence of the target DNA or RNA sequences to activity of a target protein, they can be used in the manipulation of activity of a target protein, they can be used in the manipulation of clissue e.g. tissue differentiation, both in vivo and in ex vivo tissue cultures. The method provides an efficient and rapid solid-phase method control of the the synthesis of thiourea and S-methylthiourea or alkoxy thiourea linkages in solution or on solid phase. Example 7; Fig 16; 48pp; English. of alkyl

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ; 0 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 20; Conservative Query Match

2709 AAAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAAAA 20 ч

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AAS04112 standard; DNA; 20 BP 29-AUG-2001 (first entry) AAS04112; RESULT 443 AAS04112/

Human SCF (stem cell factor) cDNA universal PCR primer 220-7. 

Human, stem cell factor, SCF, early haematopoietic progenitor cell, blood disorder, leukaemia; Hodgkin's disease, lymphoma, splenomegaly, anaemia, Kala azar, septicemia, malaria, hypopigmentation disorder, PCR primer; 88.

Homo sapiens.

US6207417-B1

27-MAR-2001

95US-00482918.

07-JUN-1995;

90US-00537198. 89US-00422383 90US-00589701 93US-00172329 11-JUN-1990; 24-AUG-1990; 01-OCT-1990; 21-DEC-1993; 16-OCT-1989;

ZSEBO K M (ZSEB/)

BOSSELMAN R A. SUGGS S V. MARTIN F H. (BOSS/) (SUGG/) (MART/)

Martin FH; Suggs SV, Zsebo KM, Bosselman RA,

WPI; 2001-298941/31.

Novel nucleic acids encoding stem cell factor useful for treating disorders involving blood cells, e.g. leukemia, splenomegaly, Hodgkin's

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The present sequence for universal PCR primer 220-7 is 1 of 8 universal oligomucleotides (AAS04110-AAS04117) used in the isolation of the human SCF (stem cell factor) DNNA sequence. The present invention relates to novel stem cell factors (AAU02453-AAU02466, AAU02461) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells concluding early haematopoietic progenitor cells. The invention also describes SCF peptides (AAU02462-AAU02481) and the oligonucleotides (AAS04081-AAS04117) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in gene therapy. It is useful for tracting disorders involving blood cells such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple compestive splenomegaly, Kala azar, sarcoidosis, military tuberculosis, disseminated fungus disease, Pulminting septicemia, malaria, vitamin B12 and folic acid deficiency, principanta deficiency, and hypopigmentation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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disease, Kala azar, anemia and septicemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorders such as piebaldism and vitiligo
                                                    Example 3; Fig 12C; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20; Conservative
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20 CTAAAAAAAAAAAAAA ВЪ AAH45787 standard; DNA; 20 (first entry) 07-SEP-2001 AAH45787; RESULT 444 AAH45787 임

2707 CTAAAAAAAAAAAAAAA 2726

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Nucleic acid amplification; adapter DNA; human; PCR primer; Human KUAPP70 gene PCR primer SEQ ID NO: 39. 16-NOV-2000; 2000WO-JP008073. WO200138572-A1. Homo sapiens. 31-MAY-2001. 

88.

99JP-00330726 25-JUL-2000; 2000JP-00224663 (TAKI ) TAKARA SHUZO CO LTD. 19-NOV-1999;

Amplifying nucleic acids with base sequences of mRNAs in sample while sustaining the ratio among them used to monitor mRNA expression, applicable in producing e.g. cRNA library and DNA microarrays.

Kato I;

Mineno J, Asada K,

Terada M,

Sasaki H,

Aoyagi K,

WPI; 2001-355947/37.

Example 1; Page 63; 67pp; Japanese.

The present invention describes a method of amplifying nucleic acids, involving forming a single-stranded DNA to an mRNA in a sample with a primer, synthesising a DNA strand complementary to the single-stranded DNA to form a double-stranded DNA, adding a single or double-stranded adapter DNA to the double-stranded DNA, and amplifying the DNA strand using a second primer with a nucleic acid sequence in the adapter DNA.

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This can be used to amplify nucleic acids to monitor mRNA expression, which is applicable in producing e.g. cRNA libraries, cDNA libraries, DNA introservays or membrane arrays in gene engineering and gene expression analysis, and in drug development and health maintenance and management. The present sequence is a PCR primer described in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated non-human mammalian stem cell factor polypeptide stimulating growth of early hematopoietic progenitor cells, useful for treating aplastic anemia, lymphoma, Letterer-Siwe disease, Kala azar, sarcoidosis.
                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; rat; mammal; stem cell factor; SCF; cell growth stimulation; gene therapy; haematopoietic disorder; aplastic anaemia; leukaemia; neurological damage; intestinal damage; infertility; AIDS; SCID; severe combined immunodeficiency; PCR primer; ss.
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                                                                                                                                                   0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalian stem cell factor PCR primer SEQ ID NO: 33.
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                                                                                                                    Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Fig 12C; 209pp; English
                                                                                                                                                                                                                        1745 CCCTCCCTGTTCGTGTACCC 1764
                                                                                                                                                                                                                                           89US-00422383.
90US-00537198.
90US-00573616.
90US-00589701.
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                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 20; Conservative
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                                                                      present se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
25-NOV-1992;
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DB 1; Length 20;

0.7%; Score 20;

Query Match

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The present sequence for universal PCR primer 220-7 is 1 of 8 universal oligonucleotides (AAH23888-AAH23895) used in the isolation of the human oligonucleotides (AAH23888-AAH23895) used in the isolation of the human SC SCF (setem cell factor) cDNA sequence. The present invention relates to novel stem cell factors (AAB73561-AAB73568, AAB73571-AAB73575) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells including early haematopoietic progenitor cells. The invention also describes SCF peptides (AAB73578-AAB73597) and the oligonucleotides (AAB73889-AAB73887) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in gene herapy. It is useful for treating disorders involving blood cells such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, anaemia, congestive splenomegaly, Rala azar, sarcoidosis, military tuberculosis, congestive splenomegaly.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New stem cell factor polypeptides and their analogs which stimulate growth of early hematopoietic progenitors, useful for treating aplastic anemia, carcinoma, multiple myeloma, vitiligo, kala azar, Hodgkin's
                                                                                                                                                                                                                                                                                                                       Human; stem cell factor; SCF; early haematopoietic progenitor cell;
blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly;
anaemia; Kala azar; septicaemia; malaria; hypopigmentation disorder;
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                                                                                                                                                                                                                                                                                     Human SCF (stem cell factor) cDNA universal PCR primer 220-7.
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100.0%; Pred. No. 6.8e+02;
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              Pred. No. 6.8e+02;
Mismatches 0;
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Matches 20; Conservative 0; Mismatches
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100.0%; Pic
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90US-00537198.
90US-00573616.
90US-00589701.
91US-00684535.
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                                                                                                                                                                                      AAH23890 standard; DNA; 20
                                                                                                                                                                                                                                                       (first entry)
               Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bosselman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-256683/26.
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                                                                                                                                                                                                                                                                                                                                                                            PCR primer; ss.
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24-AUG-1990;
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                                                                                                                                                                                                                       AAH23890;
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Human, stem cell factor; SCF; haematopoietic progenitor cell; blood disorder; Hodgkin's disease; vitamin B12; folic acid deficiency; hypopigmentation disorder; viral disorder; AIDS; PCR primer; ss.

Human stem cell factor (SCF) cDNA universal PCR primer 220-7.

(first entry)

24-OCT-2001

AAS10448;

AAS10448 standard; DNA; 20 BP.

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RESULT 448
                        AAS10448/
                                                        The present sequence for universal PCR primer 220-7 is 1 of 8 universal oligonucleotides (AAS04211-AAS04218) used in the isolation of the human SCF (stem cell factor) CDNA sequence. The present invention relates to novel stem cell factors (AAU02761-AAU02767, AAU02775-AAU02775, AAU02775) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells including early haematopoietic progenitor cells. The invention also describes SCF peptides (AAU02777-AAU02794) and the oligonucleotides.

(AAS04182-AAS04210) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in the isolation of human, and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in the isolation of human, acute leukaemia, multiple such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, anaemia, congestive sphenomegaly, Kala azar, sarcoidosis, military tuberculosis, disseminated fungus disease, Pulminating septicemia, malaria, vitamin B12 and folic acid deficiency, pyridoxine deficiency, and hypopigmentation construction as piebaldism and vitiligo
                                                                                                                                                                                                                                                                     Human, stem cell factor, SCF, early haematopoietic progenitor cell,
blood disorder, leukaemia, Hodgkin's disease, lymphoma, splenomegal,
anaemia, Kala azar, septicemia, malaria, hypopigmentation disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated DNA sequence, encoding polypeptide product useful for stimulating growth of early hematopoietic progenitor cells.
                                                                                                                                                                                                                                   Human SCF (stem cell factor) cDNA universal PCR primer 220-7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 20; DB 1; Le
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suggs SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Fig 12C; 167pp; English.
CTAAAAAAAAAAAAAA 2726
                                   CTAAAAAAAAAAAAAA 1
                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89US-00422383.
90US-00537198.
90US-00573616.
90US-00589701.
92US-00982255.
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                                                                                                                           AAS04213 standard; DNA; 20
                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-281051/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                              PCR primer, ss.
                                                                                                                                                                                                 29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                      US6218148-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-JUN-1990;
24-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                          17-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-0CT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-1990
                                                                                                                                                             AAS04213;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zsebo KM,
                                20
2707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                         447
                                                                                         RESULT
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The present sequence for universal PCR primer 220-7 is 1 of 19 PCR primers (AAS10435-AAS10453) used to amplify various portions of the human SCF CDNA sequence. The sequence is described in an invention relating to novel stem cell factors, the polynucleotides encoding them and methods for producing the stem cell factors. The methods involve increasing the number of early haematopoietic progenitor cells in human peripheral blood by administering a haematopoietically effective human stem cell factor polypeptide. The methods are useful for the treatment of blood disorders, including myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodkin's disease, a disease, leading myelofibrosis, viemnn-Pick disease, refractory anaemia, malaria, vitamin B12 and folic acid deficiency, hypopigmentation disorders i.e. piebaldism and viral induced disorders, including AIDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ...
Example 3; Fig 12C; 210pp; English.
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Best Local Similarity
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ABS77742/c
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Matches
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Gaps

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Indels

2707 CTAAAAAAAAAAAAAAA 2726

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Local Similarity 100. nes 20; Conservative

Best Loca Matches

CTAAAAAAAAAAAAAAA 1

20

Increasing the number of early hematopoietic progenitor cells in the peripheral blood useful for the treatment of blood disorders including Hodgkin's disease comprises the administration of human stem cell factor.

Suggs SV, Martin FH;

Bosselman RA,

Zsebo KM,

WPI; 2001-407312/43.

(ZSEB/) ZSEBO K M. (BOSS/) BOSSELMAN R A. (SUGG/) SUGGS S V. (MART/) MARTIN F H.

90US-00589701. 91US-00684535. 92US-00982255. 93US-00172329.

90US-00537198.

95US-00449653 89US-00422383

24-MAY-1995;

16-OCT-1989

11-JUN-1990 24-AUG-1990

01-OCT-1990

21-DEC-1993

5-NOV-1992 10-APR-1991

US6248319-B1. Homo sapiens.

19-JUN-2001.

Tue Nov

Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; conneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;

scleroderma, hypertrophic scar,

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The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lession, theumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
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                                                                                                                                                                                                                                                                                  tumour metastasis; precancerous legion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectesia, haemophilac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma hypertrophic scars The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                   Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                Angiogenesis inhibitory oligonucleotide #226.
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100.0%; Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page 23; 276pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (COLE-) COLEY PHARM GROUP INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-DEC-2001; 2001WO-US048458.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-DEC-2000; 2000US-025534P.
                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-566690/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200253141-A2.
                                                                                  13-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
ABS77742
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administering at least one antifaring instructions and molecule. Also included is a kit comprising a first container housing them to a subject nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Obler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, naemophilac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic

acid of the invention

Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.

(COLE-) COLEY PHARM GROUP INC.

WPI; 2002-566690/60.

Bratzler RL;

14-DEC-2001; 2001WO-US048458. 14-DEC-2000; 2000US-025534P.

WO200253141-A2.

Synthetic

11-JUL-2002.

The invention relates to inhibiting angiogenesis in a subject,

Claim 2; Page 29; 276pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                        Gaps
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                                                    0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.88+02; ive 0; Mismatches 0; Indels
Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Angiogenesis inhibitory oligonucleotide #560.
                                                                                                                                                                                         2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                ABS78076 standard; DNA; 20 BP.
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                                                                                                                               20; Conservative
                                                                                                 Best Local Similarity
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                                                                       Query Match
                                                                                                                                     Matches
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Gaps

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Length 20; 0; Indels

DB 1; Le

Mismatches Score 20; Pred. No.

2709 AAAAAAAAAAAAAAAAA 2728

Conservative

Local Similarity hes 20; Conserv

Matches

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Query Match

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20

ABS78072 standard; DNA; 20 BP

RESULT 450

ABS78072,

ABS78072

2X2X5X8X

Angiogenesis inhibitory oligonucleotide #556.

(first entry)

Bratzler RL;

Inhibiting

11-JUL-2002

Synthetic.

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cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the
                                                                                                                                                                                                                                                                                                                                                                                                                        cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, tissue cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, ossophageal cancer, eye cancer, Hodgkin's lymphoma, inver cancer, lung cancer, Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, pancreatic cancer, proseate cancer, rhabdomyosarcoma, skin
                                                                                                                                                                                                                                                 Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to methods for treating or preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 309; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exemplification of the invention
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                           22-JUN-2001; 2001WO-US020154.
                                                                     22-JUN-2000; 2000US-0213346P.
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                                                                                                               (IOWA ) UNIV IOWA RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20; Conservative
                                                                                                                                                             Weiner G, Hartmann G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hartmann G;
                                                                                                                                                                                                     WPI; 2002-154611/20.
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                                                                                                                                                                                                                                                                                                                    developing cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL38648;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangaectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and
                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbone; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ch 0.7%; Score 20; DB 1; Length 20; Similarity 100.0%; Pred. No. 6.8e+02; 20; Conservative 0; Mismatches 0; Indels
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/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory nucleic acid SEQ ID NO: 838.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page 29; 276pp; English.
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scleroderma; hypertrophic scar.
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ABL39402 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                 (COLE-) COLEY PHARM GROUP INC
                                                                                                                                                                                                                       14-DEC-2000; 2000US-0255534P.
                                                                                                                                                                              14-DEC-2001; 2001WO-US048458
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Best Local Similarity
Matches 20; Conserv
                                                                                     WO200253141-A2.
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modified_base
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27-DEC-2001

Synthetic

ABL39402;

RESULT 452 ABL39402/

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Gaps

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0; Indels

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The present invention relates to methods for treating or preventing developing administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, brain and central nervous system (CNS) cancer, brasse cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oesophageal cancer, elung cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, kidney cancer, larynx cancer, pancreatic cancer, plodgkin's lymphoma, melanoma, mycloma, oral carity cancer, over an earlier, orancer, cancer, testicular cancer, ribabdomyosarcoma, skin cancer, testicular cancer, and uterine cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of developing cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antibody-induced cell lysis; cancer; immunostimulatory; CD20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunostimulatory nucleic acid SEQ ID NO: 839.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. No. 6.8 cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      angiogenesis, metastasis, cytostatic, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure, Page 309, 312pp, English.
                                                                                                   Disclosure; Page 95; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2709 AAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                   exemplification of the invention
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Best Local Similarity 100.0
Matches 20, Conservative
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treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, cesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, stomach cancer, prostete cancer, rhaddomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Angiotensin-I converting enzyme; ACB; CD14; receptor; SNP; single-nucleotide polymorphism; PCR; primer; ss.
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                          Length 20;
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                                                                                                                                                                                                                                                      0.7%; Score 20; DB 1; L
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD14 receptor PCR primer SEQ ID NO 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL54775 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JUN-2002 (first entry)
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Best Local Similarity 100.C
Matches 20, Conservative
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Best Local Similarity
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ABL54775/c
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Gaps

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2709 AAAAAAAAAAAAAAAAA 2728

20; Conservative

Matches

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20 AAAAAAAAAAAAAAAA 1
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RESULT 456

ABK65035;

ABK65035 standard; DNA; 20 BP

**ABK65035** 

02-JUL-2002 (first entry)

Nanoparticle-oligonucleotide #55

Nanoparticle-oligonucleotide; nanofabrication; nucleic acid detection;

Synthetic.

WO200218643-A2

07-MAR-2002

10-AUG-2001; 2001WO-US025237

11-AUG-2000; 2000US-0224631P. 08-DEC-2000; 2000US-0254392P. 11-DEC-2000; 2000US-025335P. 12-JAN-2001; 2001US-00760500. 28-MAR-2001; 2001US-00820279.

(NANO-) NANOSPHERE INC.

Storhoff JJ, Elghanian R; Mirkin CA, Letsinger RL, Mucic RC, SI Taton TA, Garimella V, Li Z, Park S; Mirkin CA,

WPI; 2002-258024/30.

Detecting nucleic acid, useful for diagnosis of genetic, viral or bacterial disease, comprises hybridizing nanoparticles with attached oligonucleotides to nucleic acid and detecting change brought about by hybridization.

Example 18; Page 410; 412pp; English.

The invention relates to a method of detecting a nucleic acid (NA) having at least 2 portions comprising: (a) providing nanoparticles (NP) with at least 2 portions comprising: (a) providing nanoparticles (NP) with a trached oligonuclecides (OGN), where OGN has a sequence complementary to the sequence of NA; (b) contacting NP under conditions

CC effective to allow hybridisation of OGN with NA; and (c) observing a detectable change brought about by hybridisation of OGN with NA. The method is useful for detecting a nucleic acid separating a selected on uncleic acid from others and methods of nanofabrication. Detecting analytes such as nucleic acids and proteins are useful for the diagnosis of generic, bacterial and viral diseases. The OGN-NP conjugates that use cyclic disulphide linkers improve the sensitivity of diagnostic assays.

CC polic disulphide linkers improve the sensitivity of diagnostic assays.

CC plud to be approximately 10 times more sensitivity than assays employing comprising a steroid astached to a cyclic disulphide have been found to be approximately 10 times more sensitive than assays employing conjugates prepared using alkanethiols or acyclic disulphides as the directly in PCR solutions. Therefore conjugates added as probes to a DNA carried through the 30 or 40 heating coling cycles of the PCR and are still able to detect the amplicons without opening the tubes and causing contamination. ABK64981-ABK65055 represent nanoparticle-oligonucleotides of the invention 

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Gaps . 0 0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. No. Query Match
Best Local Similarity 100.
Matches 20; Conservative

2709 AAAAAAAAAAAAAAA 2728 ð g

RESULT 457 ABK65050

BP. ABK65050 standard; DNA; 20

ABK65050;

(first entry) 02-JUL-2002

Nanoparticle-oligonucleotide #70

Nanoparticle-oligonucleotide; nanofabrication; nucleic acid detection;

Synthetic.

WO200218643-A2.

07-MAR-2002.

10-AUG-2001; 2001WO-US025237

11-AUG-2000; 2000US-0224631P. 08-DEC-2000; 2000US-0254392P. 11-DEC-2000; 2000US-02535P. 12-JAN-2001; 2001US-00760500. 28-MAR-2001; 2001US-00820279.

(NANO-) NANOSPHERE INC.

Storhoff JJ, Elghanian R; Mirkin CA, Letsinger RL, Mucic RC, St Taton TA, Garimella V, Li Z, Park S;

WPI; 2002-258024/30.

Detecting nucleic acid, useful for diagnosis of genetic, viral or bacterial disease, comprises hybridizing nanoparticles with attached oligonuclectides to nucleic acid and detecting change brought about hybridization.

Example 24; Fig 44; 412pp; English.

The invention relates to a method of detecting a nucleic acid (NA) having at least 2 portions comprising: (a) providing nanoparticles (NP) with at least 2 portions comprising: (a) providing nanoparticles (NP) with a tatached oligonuclectides (OGN), where OGN has a sequence complementary to the sequence of NA; (b) contacting NA and NP under conditions ceffective to allow hybridisation of GON with NA. The detectable change brought about by hybridisation of OGN with NA. The detectable change brought about by hybridisation of OGN with NA. The conclet acid from others and methods of nanofabrication. Detecting analytes such as nucleic acids and proteins are useful for the diagnosis of genetic, bacterial and viral diseases. The OGN-NP conjugates that use cyclic disulphide linkers improve the sensitivity of diagnostic assays.

In particular assays using OGN-NP conjugates prepared using linkers of omprising a steroid residue attached to a cyclic disulphide have been found to be approximately 10 times more sensitivity than assays employing comprising a steroid residue attached to a cyclic disulphides as the directly in PCR solutions. These more sensitive than assays employing conjugates prepared using alkanethiols or acyclic disulphides as the directly in PCR solutions. Therefore conjugates and as probase to a DNA target to be PCR amplified can be carried through the 30 or 40 heating colling cycles of the PCR and are still able to detect the amplicons without opening the tubes and causing contamination. ABK64981-ABK65055 represent nanoparticle-oligonucleotides of the invention 

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Query Match

0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels

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2709 AAAAAAAAAAAAAAAAA 2728
                                 1 адададададададад 20
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RESULT 458 AAD35465, 

BP. AAD35465 standard; DNA; 20

AAD35465;

(first entry) 25-JUL-2002 Rat SCF 5' cDNA amplifying PCR primer, 220-7.

Rat; stem cell factor; SCF protein; leucopaenia; thrombocytopaenia; anaemia; myelosuppression; nerve damage; myeloproliferative disorder; infertility; neoplasa; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia; multiple myeloma; sarcoidosis; Hodgkin's disease; lymphoma; daucher's disease; Niemann-Pick disease; Letteer-Siwe disease; refractory erythroblastic anaemia; Kala azar; bi Guglielmo syndrome; congestive splenomegaly; splenic pancytopaenia; diseaminated fungus disease; Polminating septicaemia; piebaldism; AIDS; acquired immune déficiency syndrome; malaria; military tuberculosis; pyridoxine deficiency; vitemin Bl2 déficiency; folic acid deficiency; blamond Blackfan anaemia; hypopigmentation disorder; vitiligo; PCR; primer;

Rattue sp.

US2002018763-A1.

14-FEB-2002

98US-00005243 12-JAN-1998; 95US-00449653 24-MAY-1995;

(ZSEB/) ZSEBO K M.

BOSSELMAN R A. SUGGS S V. MARTIN F H. (MART/) BOSS/) SUGG/)

Suggs SV, Martin FH; Zsebo KM, Bosselman RA,

WPI; 2002-350789/38

Novel non-naturally-occurring stem cell factor polypeptide, useful for treating leucopenia, thrombocytopenia, anemia and for enhancing engraftment of bone marrow during transplantation in a mammal.

Example 3; Fig 12C; 217pp; English.

The present invention relates to novel non-naturally-occurring stem cell factor (SCF) polypeptides having an amino acid sequence sufficiently duplicative of that of naturally-occurring SCF to allow possession of haematopoietic biological activity of naturally occurring SCF. Sequences of the invention are useful for treating leucopeenia, thrombocytopaenia, anemia and for enhancing bone marrow recovery in treatment of radiation, engraftment of bone marrow during transplantation in mammals and chemical or chemotherapeutic induced bone marrow aplasia or myelosuppression. They are also useful for treating acquired immune deficiency in a human, nerve damage, neoplasia, infertility, myeloproliferative disorder, intestinal damage in a mammal. SCF sequences are useful for proparing biologically active polymer polypeptide adduct, for enhancing transfection of early active polymer polypeptide adduct, for enhancing transfection of early active polymer polypeptide adduct, for enhancing transfection of early active polymer polypeptide adduct, for enhancing transfection of agene into a mammal. They are useful for treating myelofibrosis, myeloscapetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, costeopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, cletterer-Siwe disease, refractory erythroblastic anaemia, Di Guglielmo syndrome, congestive splenomegaly, Kala azar, sarcoidosis, primary splenic pancytopaenia, disseminated fungus disease, malaria, military

pyridoxine deficiency, vitamin B12 tuberculosis, Fulminating septicaemia, pyridoxine deficiency, vitamin Bliand folic acid deficiency, Diamond Blackfan anaemia, hypopigmentation disorders such as piebaldism, AIDS (acquired immune deficiency syndrome) and vitiligo. The present sequence is a PCR primer which is used for amplifying the 5' and of rat SCF CDNA. This sequence is used in the exemplification of the invention

Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other; 8888888

ö Length 20; 0; Indels 0.7%; Score 20; DB 1; Le 100.0%; Pred. No. 6.8e+02; iive 0; Mismatches 0; 20; Conservative Local Similarity Query Match Matches

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Gaps

2707 CTAAAAAAAAAAAAAAA 2726 ઠે

CTAAAAAAAAAAAAAAAA 1 20 d

BP. ABS73849 standard; DNA; 20

ABS73849;

05-DEC-2002

(first entry)

SCF universal oligonucleotide 220-7.

Stem cell factor; SCF; blood-forming system; blood cell disorder; haematopoietic system; metastatic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; malaria; vitiligo; refractory erythroblastic anaemia; miliary tuberculosis; cytostatic; disseminated fungus disease; haematopoietic; tuberculostatic; antifungal; antimalarial; dermatological; ss. 

Synthetic.

EP1241258-A2

18-SEP-2002.

04-OCT-1990; 2002EP-00008587

90US-00537198. 89US-00422383 16-OCT-1989; 11-JUN-1990;

90WO-US005548. 24-AUG-1990;

90US-00589701. 95EP-00105391 28-SEP-1990; 01-OCT-1990; 04-OCT-1990;

04-OCT-1990

(AMGE-) AMGEN INC.

Martin FH; Æ, Bosselman Suggs SV, Zsebo KM,

WPI; 2002-684093/74.

Production of a human stem cell factor (SCF) polypeptide for treating disorders involving blood cells, such as leukemia, comprises culturing mammalian cells comprising non-human SCF promoter DNA linked to DNA encoding the human SCF

Example 3; Fig 12C; 120pp; English.

The present invention relates to novel stem cell factors (SCFs), polynucleotide sequences encoding the SCFs, and methods of producing them. SCFs are involved in the blood-forming (haematopoietic) system in mammals, particularly humans. The method of the invention is useful for the production of human SCF. The stem cell factors are useful to treat disorders involving blood cells e.g. metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, refractory erythroblastic anaemia, miliary tuberculosis, disseminated fungus disease, malaria, and vitiligo. The present sequence representing a

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ABL36232 standard; DNA; 20
                                                                     08-APR-2002
                                                ABL36232;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for the synthesis of a polynucleotide which involves coupling a second nucleoside to a first nucleoside through a phosphite linkage, where the second nucleoside has a non-carbonate protecting group protecting a hydroxyl, and exposing the product to a composition which concurrently oxidizes the phosphite formed to a phosphate and deprotects the protected hydroxyl of the second nucleoside. The method is useful for synthesizing the polynucleotides, for carrying out either 3' to 5' or 5' to 3' synthesis and for fabricating an addressable array of polynucleotides on a substrate. The present sequence is an oligonucleotide produced to demonstrate the method
                                                                                Gaps
universal oligonucleotide for SCF DNA is used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthesis of polynucleotide useful during fabrication of an array involves coupling nucleoside phosphoramidite and a solid-supported nucleoside and treating the product with an oxidation/deprotection
                                                                                                                                                                                                                                                                             Oligonucleotide synthesis; polynucleotide array; protecting group; oxidation; ss.
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Pred. No. 6.8e+02;
                                                       0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                              0; Indels
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                                  Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                      Oligonucleotide synthesis method related DNA #1.
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                                                                                                    2707 CTAAAAAAAAAAAAAAA 2726
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                                                                                                                         20 CTAAAAAAAAAAAAAAA
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                                                                                                                                                                                    AAL45122 standard; DNA; 20
                                                                                                                                                                                                                                (first entry)
                                                                                Conservative
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Best Local Similarity
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                                                                   Local Similarity
             present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dellinger DJ,
                                                                                                                                                                                                                                                                                                                                     EP1176151-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       composition.
                                                                                                                                                                                                                                24-MAY-2002
                                                                                                                                                                                                                                                                                                                                                            30-JAN-2002.
                                                                                                                                                                                                                                                                                                                Synthetic.
                                                       Query Match
                                                                                                                                                                                                         AAL45122;
                                                                   Best Loca
Matches
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                                                                                                            Skin disorder; psoriasis; atopic dermatitis; allergic contact dermatitis; alopecia areata; skin cancer; Mycobacterium vaccae; melanoma; cytostatic; antipsoriatic; dermatological; antiinflammatory; antiallergic; Th2 immune response; immunomodulatory; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method of inhibiting skin inflammation associated with a skin disorder selected from psoriasis, atopic dermatitis, and allergic contact dermatitis, which involves administering a composition containing delipidated and deglycolipidated Mycobacterium vaccae cells or M. vaccae culture filtrate. The skin disorder to be treated may also include alopecia areata, and skin cancers such as basal cell carcinoma, squamous cell carcinoma and melanoma. The composition acts by inhibiting the Th2 immune response. The present sequence is a probe described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid detection method; nanoparticle-oligonucleotide conjugate; nanoparticle; viral RNA detection; bacterial DNA detection; fungal DNA detection; nanoprobe conjugate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibiting skin inflammation associated with skin disorder e.g. psoriasis, by administering composition comprising delipidated and deglycolipidated Mycobacterium vaccae cells or Mycobacterium vaccae culture filtrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 20; DB 1; L
100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. ...
                                                    M tuberculosis rRNA probe SEQ ID NO: 83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Col 99-100; 116pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GENE-) GENESIS RES & DEV CORP LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prestidge R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AAAAAAAAAAAAAAAA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00324542.
                                                                                                                                                                                                                                                         Mycobacterium tuberculosis.
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tan PLJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-138361/18.
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                                                                                                                                                                                                                                                                                                              JS6328978-B1
                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUN-1999;
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Gaps

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2709 AAAAAAAAAAAAAAAA 2728

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Conservative

20 AAAAAAAAAAAAAAAAA 1

fungal DNA detection; nanoprobe conjugate; ss.

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                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a method of detecting (M1) a nucleic acid having two portions, involving providing nanoparticles having oligonucleotides attached to it, which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles, to allow hybridisation of oligonucleotides with two or more portions of nucleic acid, and observing a detectable change brought about by hybridisation. (M1) nanoparticles (1), nanoparticle oligonucleotide conjugates (II) and the aggregate probe are useful for detecting two or more nucleic acids (from a biological source) having at least two or more nucleic acids (from a biological source) having at least two or portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic with a disease, synthetic, or structurally-modified natural or synthetic (II) is useful for preparing a nanoprobe conjugate for detecting an analyte, and for detecting an analyte, and for detecting an encleic acid having two portions from other nucleic acids. (I), (II) and the are useful for detecting an analyte (especially analyte) and an analyte (especially analyte).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                      Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid detection method; nanoparticle-oligonucleotide conjugate; nanoparticle; viral RNA detection; bacterial DNA detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                      Mucic RC, Storhoff JJ, Elghanian R; Z, Park S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polyvalent analyte) in a sample. This sequence represents a polynucleotide used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid detection method associated polynucleotide #70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .7%; Score 20; DB 1; Length 20; 3.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                              Example 18; Page 437; 442pp; English.
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                                                                                                                                                                                                                                                        Letsinger RL, Mucio
Garimella V, Li Z,
                                                                                                                  08-DEC-2000; 2000US-0254418P.
11-DEC-2000; 2000US-0255235P.
11-DEC-2000; 2000US-0255236P.
12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-0082279.
09-APR-2001; 2001US-0282660P.
10-AUG-2001; 2001US-0282660P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS64688 standard; DNA; 20 BP
                                                                              07-DEC-2001; 2001WO-US046418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.2
Matches 20; Conservative
                                                                                                                                                                                                                               (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                                                                                WPI; 2002-608256/65.
                           WO200246472-A2
                                                                                                          08-DEC-2000;
                                                      13-JUN-2002
                                                                                                                                                                                                                                                       Mirkin CA,
  Synthetic.
                                                                                                                                                                                                                                                                      raton TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS64688;
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The invention describes a method of detecting (M1) a nucleic acid having two portions, involving providing nanoparticles having oligonucleotides attached to it, which has a sequence complementary to sequence of two cortions of nucleic acid, contacting nucleic acid and nanoparticles, to allow hybridisation of oligonucleotides with two or more portions of nucleic acid, and observing a detectable change brought about by hybridisation. (M1), nanoparticles (1), nanoparticle oligonucleotide hybridisation. (M1), nanoparticles (1), nanoparticle-oligonucleotide conjugates (II) and the aggregate probe are useful for detecting two or more nucleic acid (from a biological source) having at least two cortions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. (II) is useful for preparing a nanoprobe conjugate for detecting an analyte, and for detecting an nucleic acid having two portions from other nucleic acids. (I), (II) and conjugate probe are useful for debatication, and for separating a salected conjugate probe are useful for describing an analyte (especially conjugate), in a sample of the separating an analyte (especially conjugate).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Elghanian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Letsinger RL, Mucic RC, Storhoff JJ,
Garimella V, Li Z, Park S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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Best Local Similarity 100.0%; Pred. No. ο.σ
Αποποκοτίνο Ο; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN87103 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                  11-DEC-2000; 2000US-0255236P.
12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-00820279.
09-APR-2001; 2001US-028540P.
10-AUG-2001; 2001US-00927777.
                                                                                                                                                                                                                                                                                                               2000US-0254418P.
2000US-0255235P.
                                                                                                                                                                                                        07-DEC-2001; 2001WO-US046418
                                                                                                                                                                                                                                                                                  2000US-0254392P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-608256/65.
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                                                                      WO200246472-A2
                                                                                                                                                                                                                                                                                                                   08-DEC-2000;
                                                                                                                                                                                                                                                                                  08-DEC-2000;
                                                                                                                                            13-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mirkin CA,
Taton TA,
Synthetic
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Unidentified.

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The present invention describes a non-antibody protein, comprising a domain having an immunoglobulin-like fold, derived from a reference brotein having a mutated amino acid sequence, where the non-antibody protein brinds with a Kd at least as tight as 10 nM to a compound that is not bound as tightly by the reference protein. The non-antibody protein is useful as scaffcilds for selecting or designing a protein framework with specific and favourable properties, e.g. for binding any antigen of interest, or for destroying or inactivating antibody molecules. The non-antibody protein is also useful in all areas where antibodies are used, e.g. research, therapeutic or diagnostic fields, and for screening novel binding proteins useful in the above-mentioned fields. The present proteins have thermodynamic properties superior to those of natural antibody mimics exhibit improved biophysical properties, such as antibody mimics exhibit improved biophysical properties. Such as concentrations. In addition, these molecules are readily expressed and folded in prokaryotic systems (e.g. yeast), or in in vitro translation systems (e.g. rebabit reticulocyte lysate system). Furthermore, these proteins are extremely amenable to affinity maturation techniques involving multiple cycles of selection, e.g. in vitro selection using RNA-protein fusion technology, phage display or yeast display systems. The present sequence is used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                 New non-antibody proteins having an immunoglobulin fold, useful in research, therapeutic or diagnostic fields, particularly as scaffolds for designing proteins with specific properties, e.g. for binding any antigen
Protein scaffold; antibody; binding protein; immunoglobulin;
tumour necrosis factor alpha; TNF-alpha; protein framework; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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100.0%; Pred. No. 6.8e+02;
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Best Local Similarity 100.0%; Pred. No. 6.8
Matches 20; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                               Kuimelis RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 58; 94pp; English.
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                                                                                                                                                                                                         16-OCT-2001; 2001WO-US032233
                                                                                                                                                                                                                                                   16-OCT-2000; 2000US-00688566
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                                                                                                                                                                                                                                                                                                                                          Lipovsek D, Wagner RW,
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-444238/47.
                                                                                                                                                                                                                                                                                                (PHYL-) PHYLOS INC.
                                                                                                              WO200232925-A2.
                                                                                                                                                             25-APR-2002
                                                                      Synthetic
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The invention relates to a method of detecting a nucleic acid having two portions. The method involves providing nanoparticles having to oligonucleotides attached to it which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles to allow hybridisation of oligonucleotides with two or more portions of nucleic acid and observing a detectable change brought about by hybridisation. The method and aggrague probes are useful for detecting two or more nucleic acids (from a biological source) having at least two portions such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic or structurally modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. The invention is useful for preparing a nanoprobe conjugate for detecting an analyte and for detecting a nucleic acid bound to an electrode surface. It is also useful for fabrication and for separating a selected nucleic acid having two portions from other nucleic acids. The present sequence is an oligo used to illustrate the method of
                                                                                                                                                                                                                                                                                                                Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Potato; plant; mitochondrial carrier protein; elongation factor EF-2; transfertin binding protein; receptor-like protein kinase; helicase; non-long terminal repeat retroelement reverse transcriptase; overwatering; transgenic; reverse transcriptase; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred. ...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                         Example 18; Page 179; 467pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                        Park S, Taton TA, Mirkin CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABZ59815 standard; RNA; 20 BP.
                                                                                                                                             09-OCT-2001; 2001US-0327864P. 07-DEC-2001; 2001US-00008978.
                                                                                                         08-OCT-2002; 2002WO-US032088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Potato gene PCR primer dT20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20; Conservative
                                                                                                                                                                                                   (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                                                            WPI; 2003-430409/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                    WO2003035829-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DE10114063-A1
                                                                        01-MAY-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                     The invention relates to DNA sequences (I) that encode six specific plant proteins: (i) a protein (ABP60425) with mitochondrial carrier protein activity (IIa); (ii) a protein (ABP60426) with transferrin binding protein activity (IIb); (iii) a protein (ABP60427) with receptor-like protein kinase activity (IId); (v) a protein (ABP60429) with helongation factor EF-2 activity (IId); (v) a protein (ABP60429) with helongation factor EF-2 activity (IId); (v) a protein (ABP60429) with non-long creminal repeat retroelement reverse transcriptase activity (IIe); or (vi) a protein (ABP60429) with helicase activity (IIf); (I), also related sequences, derived ribozymes and antisense sequences, expression vectors, encoded proteins and antibodies against the proteins are used to produce plants with altered properties, including tolerance of overwatering. The antibodies are also used for isolation of the proteins and in immunoassays. Also (I) or their primer or probe fragments are used to screen for terminators and constitutively, aerobically or anaerobically inducible plant promoters, specifically for use in potatoes and the sequence that encodes (IId) is used to alter the translation profile in plants or since (I) are derived from potato, their promoters and the control endogenous genes. The present sequence is that of a PKR primer control endogenous genes. The present sequence is that of a PKR primer control or and a primer of the present sequence is that of a PKR primer control or and a primer of the present sequence is that of a pKR primer control or and a primer or a probe of the present sequence is that of a pKR primer control or and a primer or a probe of the present sequence is that of a primer or and a primer or and a primer or a primer or and a prime
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                                                                                                                                                                      New DNA sequences from potato, useful for producing plants with altered properties, e.g. tolerance of flooding, also related proteins, antibodies and inhibitory sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nanoparticle; 88; nucleic acid detection; viral disease; probe; human immunodeficiency virus infection; hepatitis virus infection; herpes virus infection; cycomegalovirus infection; forensic science; Epstein-Barr virus infection; bacterial disease; gene therapy; escually transmitted disease; inherited disease; pNs sequencing; paternity testing; cell line authentication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              first strand synthesis of cDNAs derived from Potato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Le
. 6.8e+02;
                                                                  (MPBC-) MPB COLOGNE GMBH MOLECULAR PLANT & PROTE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 20; DB
100.0%; Pred. No. 6.8
tive 0; Mismatches
                                                                                                      Haussuehl K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thio-modified 20dA oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                Example 1; Page 8; 26pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX79181 standard; DNA; 20 BP
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   22-MAR-2001; 2001DE-01014063
                                  22-MAR-2001; 2001DE-01014063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                      Tscharntke M,
                                                                                                                                         WPI; 2003-041808/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JS2002155462-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   used in the
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                                                                                                          Buelow L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABX79181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Matches
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The invention relates to detecting a nucleic acid (NA) having at least 2 portions, comprises providing a type of nanoparticles (NP) having attached to oligonuclectides (O) ((0) on each NP has a sequence complementary to sequence of at least 2 portions of NA), contacting NA and NP to allow hybridisation of (0) on NP with Contacting NA, contacting a detectable change brought about by hybridisation of (0) on NP with NA. The nanoparticle is useful for separating a selected nucleic acid having at least 2 portions, from other nucleic acids, and for detecting nucleic acids having at least 2 portions. The method of cusing NP is useful for detecting any type of nucleic acids which may be cusing NP is useful for detecting nucleic acids of nucleic acids of preferably, the method is useful for detecting nucleic acids for diagnosis and/or monitoring of viral diseases (human immunodeficiency virus), hepatitis virus, herpes virus, cytomegalovirus and Epstein-Barr virus), bacterial diseases, sexually transmitted diseases, inherited diseases, in forensics, in DNA sequencing, for paternity testing, for cell line authentication and for monitoring gene therapy. The method is useful in research and analytical laboratories in DNA sequencing and in the field to detect the presence of specific pathogens. Detecting nucleic acids based on observing a colour change with the maked eye is cheap, calse, simple and robust, and do not require specialised expensive equipment. The present sequence is a nanoparticle (e.g. gold particles)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nonoparticle; nucleic acid detection; hybridisation; diagnosis; sequencing; viral infection; human immunodeficiency virus; HIV; hepatitis virus; herpes virus; cytomegalovirus; Epstein-Barr virus; bacterial infection; sexually transmitted disease; inherited disorder; forensic; paternity testing; cell line authentication; gene therapy; 68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting nucleic acids having at least 2 portions comprises use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the nucleic acid sequence.
                                                                                                                                                                                                                                                                                                                        Elghanian R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                        Mucic RC, Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nanoparticle-associated oligonucleotide SEQ ID 55.
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                                 97WO-US012783.
99US-00240755.
99US-00344667.
                                                                                                                                       26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
96US-0031809P
                                                                                                                                                                                                                                                                                                                               Letsinger RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                    (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-198491/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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                                                                                                         25-JUN-1999;
26-APR-2000;
                                 21-JUL-1997
                                                                                                                                                                                                                                                                                                                           Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                             Taton TA;
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This invention describes a novel method of detecting nucleic acid having at least two portions. The method involves providing nanoparticles attached to oligonucleotides, where the oligonucleotide on each nanoparticle have a sequence complementary to a sequence of at least two portions of nucleic acid, contacting nucleic acid and nanoparticle with two complements acid. The method is useful for separating a detectable change brought about by hybridisation of the oligonucleotide nanoparticle with two or more portions of nucleic acid and observing a selected nucleic acid having at least two portions, from other nucleic acids and for detecting nucleic acids having at least two portions. The method is useful for detecting any type of nucleic acids which may be used for diagnosis of disease and in sequencing of nucleic acids. Preferably, the method is useful for detecting any type of nucleic acids for diagnosis and/or monitoring of viral infections (human immunodeficiency virus (HIV), hepatitis virus, there is virus, cytomegalovirus and Epstein-Barr virus), bacterial chiesease, in DNA sequencing, for paternity testing, for cell line concensics, in DNA sequencing, for paternity testing, for cell line casearch and analytical laboratories in DNA sequencing in the field to detect the presence of specific pathogens. Detecting nucleic acids based on observing a colour change with the naked eye is cheap, fast, simple and robust and does not require specialised expensive equipment. ABX92123 captod of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                      nse of
                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting nucleic acids having at least two portions involves use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the nucleic acid sequence.
                                                                                                                                                                                                                                                                                                                                Storhoff JJ, Elghanian R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                            Mucic RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 59; 130pp; English.
                                                                                                                                       96US-0031809P.
97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0200161P.
                                                                                                 28-SEP-2001; 2001US-00967409
                                                                                                                                                                                                                        26-APR-2000; 2000US-0200161P. 26-JUN-2000; 2000US-00603830.
                                                                                                                                                                                                                                                                                                                              Letsinger RL,
                                                                                                                                                                                                                                                                                     (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-182627/18.
                   US2002155458-A1.
                                                                                                                                       29-JUL-1996;
                                                                                                                                                                                  29-JAN-1999;
25-JUN-1999;
                                                                                                                                                               21-JUL-1997
                                                         24-OCT-2002
                                                                                                                                                                                                                                                                                                                            Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                 raton TA;
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Gaps
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0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02;
                         0; Indels
       100.0%; Pred. ...
           Best Local Similarity 100.
Matches 20; Conservative
Query Match
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2709 AAAAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAA 20

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ACD27255 standard; DNA; 20 BP ACD27255 RESULT 469 8X4X5X8

15-OCT-2003 (first entry)

Nanotechnology nucleic acid detection method associated #54.

virus detection; human immunodeficiency virus; HIV; hepatitis; herpes; cytomegalovirus; Epstein-Barr virus; bacterial disease; DNA sequencing; sexually transmitted disease; inherited disorder; forensic; Mucic RC, Storhoff JJ, Elghanian R; Nanotechnology; ss; nucleic acid detection; nanoparticle; /\*tag= a /mod\_base= OTHER /note= "OTHER= Thiol modified" " paternity testing; cell line authentication. Location/Qualifiers 21-JUL-1997; 97WO-US012783. 25-JAN-1999; 99US-00240755. 25-JUN-1999; 99US-00344667. 26-APR-2000; 2000US-020161P. 96US-0031809P 11-OCT-2001; 2001US-00975062 26-JUN-2000; 2000US-00603830 Mirkin CA, Letsinger RL, (NANO-) NANOSPHERE INC. /mod WPI; 2003-228114/22. US2002155459-A1. modified\_base 29-JUL-1996; 24-OCT-2002 Synthetic Taton TA; 

ဌ Detecting nucleic acids having 2 portions e.g. for detecting disease, comprises use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the nucleic acid sequence.

Example 18; Page 43; 129pp; English.

This invention relates to a novel method for detecting a nucleic acid having 2 portions. The method comprises providing nanoparticles having 2 portions. The method comprises providing nanoparticles to algonucleotide on each nanoparticle has a sequence complementary to a sequence of 2 portions of nucleic acid and nanoparticle are contacted to allow hybridisation of the oligonucleotide on the nanoparticle with two or more portions of cut plopanicleotide on the nanoparticle with two or more portions of or nucleic acid and observing a detectable change brought about by the hybridisation. The method of the invention is useful for separating a content nucleic acids, and content nucleic acids having 2 portions. The method of the invention is useful for detecting any type of nucleic acids which may be used for disease and in sequencing of nucleic acids for disponsis and/or monitoring of viral diseases (human immunodeficiency virus, hepters virus, herpes virus, cytomegalovirus and Epstein-Barr virus), bacterial diseases, sexually transmitted disease, inherited diseases, in forensics, in DNA sequencing, for paternity testing, for cell line authentication, for monitoring gene therapy, etc. This method involves detecting nucleic acids based on observing a colour change with the naked eye so is cheap, fast, simple and robust, and does not require content in a sequence represents a thiol specialised expensive equipment. The present sequence represents a thiol modified oligonucleotide sequence used to demonstrate the method of the invention

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Ouery Match

0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels

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2709 AAAAAAAAAAAAAAAA 2728 

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Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change,
                                                                                                                                                                                                                                                                                           Nanotechnology, nucleic acid detection, nanoparticle, ss, forensic; DNA sequencing; paternity testing; cell line authentication.
                                                                                                                                                                                                                                             Nanotechnology nucleic acid detection method oligonucleotide #54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Storhoff JJ, Elghanian R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'note= "OTHER= Thiol modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mucic RC,
                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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/mod_base= OTHER
1 AAAAAAAAAAAAAAAA 20
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99US-00240755.
99US-00344667.
                                                                                                              BP.
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25-JUN-1999; 99US-0034667.
26-AFR-2000; 2000US-020161P.
26-JUN-2000; 2000US-00603830.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-SEP-2001; 2001US-00966312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-0031809P
                                                                                                                ACD27125 standard; DNA; 20
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                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                           modified base
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21-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                          Synthetic
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                                                                                                                                                              ACD27125;
                                                                     RESULT 470
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This invention relates to a novel method for detecting nucleic acid
sequences having two portions. The method involves providing
contentials having on oligonucleotides attached to them, which has a
sequence complementary to sequence of two portions of nucleic acid,
contacting nucleic acid and nanoparticles, to allow hybridisation of
contacting nucleic acid and nanoparticles, to allow hybridisation of
contacting nucleic acid and nanoparticles, to allow hybridisation of
a detectable change brought about by hybridisation. The method of the
contacting acids (from a biological source) having at least two portions,
such as viral RNA or DNA, bacterial or fungal DNA, a gene associated with
a disease, synthetic, or structurally modified natural or synthetic RNA
a disease, synthetic, or structurally modified natural or synthetic RNA
cor DNA, or a product of a polymerase chain reaction amplification.
Nanoparticles and nanoparticle oligonuclectide conjugates of the
invention are useful for nanofabrication, and for separating a selected
invention is useful in forensics, DNA sequencing, for paternity
ctesting, cell line authentication, and monitoring gene therapy.
Ctesting, call line authentication, and monitoring gene therapy.
Ctesting, call line authentication, and monitoring gene therapy.
Ctesting, call line authentication, and monitoring gene conjugates
Cf the invention improve the sensitivity of the nucleic acid detection
assay. The present sequence represents a thiol modified oligonucleotide
cross sequence used to demonstrate the method of the invention
                                                                                                                      Example 18; Page 44; 130pp; English.
useful in forensics.
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This invention relates to a novel method for detecting a nucleic acid by contacting a nucleic acid with at least two types of nanoparticles having oligonucleotides attached, allowing hybridisation of the oligonucleotides content and observing a detectable change. The coligonucleotides on each nanoparticle have a sequence complementary to its respective portion of the sequence of the nucleic acid to be content of the invention may be used for the detection of a detected. The method of the invention may be used for the detection of a sequencing, in the field to detect the presence of specific pathogens, in the doctor's office for quick identification of an infection to assist in the correction of a nucleic acid used in, esquencing. The method of the invention detects in expensive first-line screening. The method of the invention detects nucleic acids based on observing a colour change with the naked eye. This cancel to the present sequence represents a thiol modified expensive equipment. The present sequence represents a thiol modified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of nucleic acid for, e.g. research and analytical laboratories in deoxyribonucleic acid sequencing, involves contacting nucleic acid with nanoparticles having oligonucleotides.
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Elghanian R;
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                                                                                                                                                                                                                                                                               Nanotechnology nucleic acid detection method associated #54.
                                                                                                                                                                                                                                                                                                             Nanoparticle; 88; nucleic acid detection; DNA sequencing;
           0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mucic RC, Storhoff JJ,
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/note= "OTHER= Thiol modified" "
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                                                                            2709 AAAAAAAAAAAAAAAAA 2728
                                                                                                         1 AAAAAAAAAAAAAAAA 20
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99US-00240755.
99US-00344667.
2000US-0200161P.
                                                                                                                                                                                       ACD27385 standard; DNA; 20 BP.
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Mirkin CA, Letsinger RL,
Taton TA;
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                                                                                                                                                                                                                                                     15-OCT-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                   pathogen detection.
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modified_base
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26-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                             RESULT 471
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 oligonucleotide sequence used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection of nucleic acid for, e.g. research and analytical laboratories in deoxyribonucleic acid sequencing, involves contacting nucleic acid with nanoparticles having oligonucleotides.
                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Elghanian R;
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                                                                                                                                                                                                                                                                                                                                Nanotechnology nucleic acid detection method associated #54.
                                                            0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                             Nanoparticle; 88; nucleic acid detection; DNA sequencing
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                              Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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/note= "OTHER= Thiol modified"
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                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                         2709 AAAAAAAAAAAAAAA 2728
                                                                                                                                             BP.
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99US-00240755.
99US-00344667.
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2000US-00603830.
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                                                                         Local Similaricy
nes 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                           *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
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29-JAN-1999;
25-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of nucleic acid having -2 portions used to prepare biomaterials and in nanofabrication methods, comprises providing nanoparticles, contacting nucleic acid and nanoparticles, and observing change.
                                                                                                                                                                                                                                                                                                                                                                                                                         Nanotechnology; nucleic acid detection; nanofabrication; nanoprobe; ss.
equipment, and little or no instrumentation is required. The present sequence represents a thiol modified oligonucleotide sequence used to demonstrate the method of the invention
                                                                                                                                    Gaps
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                                                                                                0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "OTHER= Thiol modified"
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Taton TA;
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                                                                                                                                                                                         BP.
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99US-00240755.
99US-00344667.
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2000US-00603830
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                                                                                                                                   20; Conservative
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                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
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25-JUN-1999;
26-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                       ACD27060;
                                                                                                 Query Match
                                                                                                                     Local
                                                                                                                     Best Loca
Matches
                                                                                                                                                                                                                                                    RESULT 473
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or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. Nanoparticle-oligonucleotide conjugates of the invention care useful for preparing a nanoprobe conjugate for detecting an analyte, and for detecting a nucleic acid bound to an electrode surface.

Nanoparticles and nanoparticle conjugates of the invention are useful for nanofabrication and for separating a selected nucleic acid having two portions from other nucleic acids. Diagnostic assays employing nanoparticle-oligonucleotide conjugates improve the sensitivity of nucleic acid detection methods and can be used to detect nucleic acids that are present in only small amounts in a sample. The invention also provides highly desirable nanoparticle-oligonucleotide conjugates. These conjugates are stable with tailored hybridisation abilities. The present conjugates at thiol modified oligonucleotide sequence used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nanotechnology; nucleic acid detection; nanofabrication; nanoprobe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detection of nucleic acid having at least two portions, by contacting nucleic acid and nanoparticles under conditions, which allows hybridization of oligonucleotides on nanoparticles with at least two portions of nucleic acid.
                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                            Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "OTHER= Thiol modified" "
                                                                                                                                                                                                                                                demonstrate the method of the invention
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29-JAN-1999; 99US-00240755.
25-JUN-1999; 99US-00344667.
26-APR-2000; 2000US-020016IP.
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modified_base
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Taton TA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 474
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Ting invention relates to a novel method for defecting nucleic acids. The method comprises providing nanoparticles with oligonucleotides attached to them, which have a sequence complementary to a sequence of two to them, which have a sequence complementary to a sequence of two to them, which have a sequence complementary to a sequence of two complementary and an anoparticles of the nucleic acid, and observing a detectable change brought about by the nucleic acid, and observing a detectable change brought about by the hypridisation. The nucleic acid to be detected must have at least two portions and the distances between these are chosen so that when the nanoparticle-oligonucleotide conjugate binds the target sequence a detectable change occurs. The method of the invention is useful for detecting two or more nucleic acids (from a biological source) having at least two portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction are useful for preparing a nanoprobe conjugates of the invention are useful for preparing a nanoprobe conjugates of the invention are useful for preparing a nanoprobe conjugates of the invention are useful for nanoparticle-oligonucleotide conjugates of the invention are useful for nanoparticle-oligonucleotide conjugates improve the sensitivity of nanofabrication methods and can be used to detect nucleic acids detection methods and can be used to detect nucleic acids or provides highly desirable nanoparticle-oligonucleotide conjugates improve the sensitivity of conjugates are stable with tailored hybridisation abilities. The present conjugates are stable with ainvention abilities. The present conjugates the method of the invention abilities. The present conjugates are invention and inven
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                This invention relates to a novel method for detecting nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             demonstrate the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KRIE/) KRIEG A M. (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003050268-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                         The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermaitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                eczema,
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                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine, non-allergic inflammatory disease, psoriasis, eczema, allergic contact dermatitis, latex dermatitis, inflammatory bowel disease, ulcerative colitis, Crohn's disease, ss.
allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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                                                         Disclosure; Page 23; 229pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory nucleic acid #533.
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Matches 20; Conservative
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                                 Gaps
                                                                                                                                                                                                                                                                                       Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine, non-allergic inflammatory disease, psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease, ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating non-allergic inflammatory diseases, such as psoriasis, ecz. allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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   Length 20;
                               Indels
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0.7%; Score 20; DB 1; Le
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0;
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Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                            Immunostimulatory nucleic acid #218.
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                                                            2709 AAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                  ACD99532 standard; DNA; 20
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                               Conservative
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               1 Similarity
20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (KRIE/) KRIEG A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                              US2003050268-A1.
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                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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Query Match
Best Local S:
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hairpin sensor useful for detecting a target nucleotide sequence in a sample, comprises a hairpin loop assembly including a complementary probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /bound_moiety= "Hairpin oligonucleotide #2" /note= "Forms a doube-stranded region with the hairpin oligonucleotide shown in examples 3, 4 and 5"
                                                                                                                                                          Hairpin sensor; hairpin loop; complementary probe; inverse repeat arm quenchable fluorescing agent; microarray; semiconductor; nanocrystal; rhodamine B-labelled dye; detection; gold support; ss.
                                                                          Hairpin target sequence, #2, used in an example of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and a quenchable fluorescing agent.
                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUN-2001; 2001US-0299460P
06-NOV-2003 (first entry)
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(LOCA/) LOCASCIO M.
(LAND/) LANDRY D P.
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misc binding
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The invention relates to detecting a nucleic acid (NA) having at least 2 portions comprises providing a type of nanoparticles (NP. e.g. colloidal gold) having oligonucleotides (O) attached (where (O) on each NP has a sequence complementary to sequence of at least two portions of NA), contacting NA, and observing a detectable change brought about by opticians of (NA, and observing a detectable change brought about by obticiantion of (O) on NP with NA. Also included are aggregate probes, over probes, substrate having NP attached to it, a metallic or semiconductor NP having (O) attached to it, inanomaterials/manostructures comprising nanoparticles and methods of nanofabrication utilising nanoparticles and satellice probes. The methods, probes nucleic acids, nanoparticles and oligonucleotides are useful for separating selected cultiplication conductor and lagonucleotides are useful for separating a selected cultiplication conductor acids, and conduction nucleic acids, and for detecting nucleic acids having at least two portions, from other nucleic acids, and for detecting and least two portions, for detecting any naving at least two portions is useful for detecting any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nanostructure; viral disease; human immunodeficiency virus infection; hepatitis virus infection; herpes virus infection; cytomegalovirus virus infection; Epstein-Barr virus; bacterial disease; sexually transmitted disease; inherited disorders; paternity testing;
                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting nucleic acids having at least 2 portions comprises use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the target nucleic acid sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; nanoparticle; colloidal gold; semiconductor; nanomaterial
                                                                               Score 20; DB 1; Length 20; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Nanoparticle labelled oligonucleotides, spacer DNA #2.
                                           Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
sequence, #2, used in an example of the invention
                                                                                      Match Local Similarity 100.0%; Pred. No. o.u.
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97WO-US012783.
99US-00240755.
99US-00344667.
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26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
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29-JAN-1999;
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type of nucleic acids which may be used for diagnosis of disease and in detecting of nucleic acids. Preferably, the method is useful for detecting nucleic acids for diagnosis and/or monitoring of viral diseases (human immunodeficiency virus, hepatitis virus, herpes virus, sexually transmitted diseases, inherited diseases, in forensics, in DNA sequencing, for paternity testing, for cell line authentication, for monitoring gene therapy, etc. The method is useful in research and analytical laboratories in DNA sequencing, in the field to detect the presence of specific pathogens, etc. Detecting nucleic acids based on observing a colour change with the naked eye is cheap, fast, simple and robust, and do not require specialised expensive equipment. The present sequence is a spacer oligonucleotide used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                         invention.
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Gaps

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0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels

20; Conservative

Nanotechnology; nucleic acid detection; nanofabrication; nanoprobe; ss. Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change. Nanotechnology nucleic acid detection method oligonucleotide #54 Elghanian R; Storhoff JJ, /mod\_base= OTHER /note= "OTHER= Thiol modified" Mucic RC, Location/Qualifiers 20 AAAAAAAAAAAAAAAAA ACD26995 standard; DNA; 20 BP. 96US-0031809P 97WO-US012783. 99US-00240755. 20-SEP-2001; 2001US-00957318 99US-00344667 2000US-0200161P 26-JUN-2000; 2000US-00603830 15-OCT-2003 (first entry) \*tag= a Letsinger RL, (NANO-) NANOSPHERE INC. WPI; 2003-615795/58. US2003049630-A1 Key modified\_base 26-APR-2000; 29-JUL-1996; 25-JUN-1999; 21-JUL-1997; 29-JAN-1999; 13-MAR-2003 Mirkin CA, Taton TA; Synthetic. ACD26995; 

Example 18; Page 43; 129pp; English

to them, which have a sequence complementary to a sequence of two controls and contacting the nucleic acid, and observing the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with two or more portions of the nucleic acid, and observing a detectable change brought about by the nucleic acid, and observing a detectable change brought about by the hybridisation. The nucleic acid to be detected must have at least two portions and the diseances between these are chosen so that when the nanoparticle-oligonuclectide conjugate binds the target sequence a detectable change occurs. The method of the invention is useful for detecting two or more nucleic acids (from a biological source) having at cleast two portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction and for preparing a nanoprobe conjugate for detecting an analyte, and for detecting a nucleic acid bound to an electrode surface. Nanoparticles and nanoparable conjugates of the invention are useful for nanofabrication and for separating a selected nucleic acid having two cortions from other nucleic acids. Diagnostic assays employing consorparticle-oligonucleotide conjugates improve the sensitivity of mucleic acid detection methods and can be used to detect nucleic acids that are present in only small amounts in a sample. The present sequence the are present at the invention of the invention and constrate the are present and encounted and can be used to detect nucleic acids that are present in only small amounts in a sample. The present sequence the are present and the present and the detection methods and can be used to detect nucleic acids that are present sequence the are present sequence. method comprises providing nanoparticles with oligonucleotides attached Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament. The invention relates to a method of treating or preventing allergy or Gaps ö ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory. Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 100.0%; Prec. ... Immunostimulatory nucleic acid #547. Disclosure; Page 13; 221pp; English 2709 AAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAA 20 뮵 02-FEB-2001; 2001US-00776479. 03-FEB-2000; 2000US-0179991P. the method of the invention ADB36933 standard; DNA; 20 Petersen DM, Query Match
Best Local Similarity 100.0 (first entry) (BRAT/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y. WPI; 2003-657977/62. US2003087848-A1. Bratzler RL, 04-DEC-2003 08-MAY-2003. Synthetic. ADB36933; RESULT 481 ADB36933 쉽 ð ö This invention relates to a novel method for detecting nucleic acids. The

BP.

(first entry)

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ADB36929 standard; DNA; 20
                                                                                        04-DEC-2003
                                              ADB36929
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asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosling and/or treating asthma and allergy especially in a hypo-responative subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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100.0%; Pred. No. 6.8e+02;
vative 0; Mismatches 0; Indels
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                                                                                                                                                                       Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches
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(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
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nes 20; Conserv
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Matches
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RESULT 483 ADB36929/c

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PCR primer for amplifying the ATP dependant DNA helicase gene #SEQ ID 68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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                                  ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
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100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indels
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Immunostimulatory nucleic acid #543.
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Best Local Similarity 100.0
"**rhes 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC24378 standard; DNA;
                                                                                                                                                                                                                                                                                                           (BRAT/) BRATZLER R L.
(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
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Page 233

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haematopoietic activity and the polynucleotides encoding them. The polypeptides are used for treating infertility, intestinal damage, myeloproliferative disorders, leucopenia, thrombocytopenia or anaemia, for improving engraftemen of bone marrow transplants, for enhancing bone marrow recovery after radiotherapy or chemotherapy and in treatment of immune deficiency, neoplasia, nerve damage, osteoporosis, metastatic carcinoma, leukaemia and miliary tuberculosis. The SCF polypeptides are also used to expand haematopoietic progenitor cells for transplantation and to prepare such cells for transfection with a gene. The SCF polypeptides and also as probes for mapping of the SCF gene, for identifying SCF-related diseases and as a marker for neighbouring genes. Antibodies raised against the polypeptides are useful in diagnosis and to remove SCF from blood. This sequence represents SCF related DNA of the
                                        New non-natural stem cell factor, useful for treating e.g. leucopenia or immune deficiency, also related nucleic acid and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of nucleic acid having at least two portions comprises contacting the nucleic acid and nanoparticles under conditions to allow hybridization of the oligonucleotides, and observing detectable change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                    invention relates to stem cell factor (SCF) polypeptides with
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Pred. No. 6.8e+02;
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100.0%; Pred. No. e...
                                                                                            Disclosure, SEQ ID NO 33; 217pp; English.
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99US-00240755.
99US-00344667.
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Taton TA;
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20; Conservative
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       WPI; 2003-851459/79.
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25-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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                                                                                                                                                                                                                                      The invention relates to a method for the amplification of DNA that maintains genes and copy number of the sequence. This method is useful for easy and operable amplification of DNA. The method was carried out by fragmentation genomic DNA, preparation of blunt end of the fragmented DNA, ligation of an adapter to the blunted DNA, PCR of the ligated DNA in 2 steps, and confirmation of the amplified APC gene. The current sequence represents a PCR primer used in an example from the invention.
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                                                                                                                                                 number of the sequence
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0
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                                                                                                                                              Amplification of DNA maintaining genes and copy number on a genome, and their ratios in the resultant DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 1 A; 10 C; 3 G; 6 T; 0 U; 0 Other;
                                (TAKA-) TAKARA BIO KK.
(KOKU-) KOKURITSU GAN CENT SOCHO.
(IYAK-) IYAKUHIN FUKUSAYO HIGAI KYUSAI KENKYU SH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Martin FH;
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                                                                                                                                                                                                   Example 5; SEQ ID NO 68; 33pp; Japanese.
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90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
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08-MAY-2001; 2001JP-00137858.
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Les 20, Conservative
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BOSSELMAN R A.
SUGGS S V.
                                                                                                           WPI; 2003-460878/44.
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24-MAY-1995
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(SUGG/)
(MART/)
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RESULT 488
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                                                                                                                                                                                             AAD64709
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                                       The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contecting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                             0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                           Nanotechnology nucleic acid detection method associated
                                                                                                                                                                                                                                                                                                                                                                                                                 Linking oligonucleotide; ss; nucleic acid detection;
nanoparticle-oligonucleotide conjugate.
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                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                    0; Mismatches
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                      SEQ ID NO 55; 129pp; English
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21-JUL-1997; 97WO-US012783.
29-JAN-1999; 99US-00344667.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-0200161P.
                                                                                                                                                                                                                                                                                                                         BP
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                                                                                                                                                                                                         Best Local Similarity 100.
Matches 20; Conservative
brought by hybridization
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                       Example 18;
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sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                               0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "Labelled with thiol group"
                                                                                                                                       Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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97WO-US012783.
99US-00240755.
99US-00344667.
2000US-00603830.
2001US-00961949.
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                                                                                                                                                                                                                                          20; Conservative
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Matches 20; Conserv
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25-JUN-1999;
26-JUN-2000; 2
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21-JUL-1997;
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Tue Nov

non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC; 88;

WO2003027328-A2.

Synthetic

probe.

Non-nucleotide probe of the invention #12.

(first entry)

25-MAR-2004

ADH59608;

ADH59608 standard; DNA; 20

ADH59608/

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of nucleic acid useful for, e.g. research and analytical laboratories in deoxyribonucleic acid sequencing, comprises contacting nucleic acid with at least two types of nanoparticles attached with oligonucleotides.
                                               Gaps
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                                                                                                                                                                                                                                                                     #54.
                  Length 20;
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                                                                                                                                                                                                                                                                                           Linking oligonucleotide; ss; nucleic acid detection;
nanoparticle-oligonucleotide conjugate.
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                DB 1; Le
6.8e+02;
                                             Mismatches
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                                                                         2709 AAAAAAAAAAAAAAAA 2728
                                                                                        Query Match 0.7%; Soc
Best Local Similarity 100.0%; PJ
                                                                                                                                                                          ADF65590 standard; DNA; 20 BP
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97WO-US012783.
99US-00240755.
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                                                                                                                                                                                                                                                                                                                                         Synthetic.
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The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic acid probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an assay for determining target genomic on comprises contacting the sample with the mixture of probes (preferably comprising 5-50 probes), contacting the sample. The method comprises contacting the sample with the mixture of probes greated by the care of the sample is an addetermining the target genomic contacting the sample by the care of the sample is an endesic or mucleic acid of the sample is an metaphase spreads, interphase nucleic acid of the sample. The genomic nucleic acid is contained in a fixed tissue or a cell, and the sample is metaphase spreads, interphase nucleic or nucleic found in paraffin embedded tissue material or frozen tissue sections. The probe is also useful in comparing a sample of genomic nucleic acid with the method comprises treating a sample of genomic nucleic acid with the method comprises treating a sample of genomic nucleic acid and control genomic nucleic acid, which are differentially labelled, the carray or both the singuals from the differentially labelled, the carray or both the singuals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus contacting under suitable hybridization conditions, contacting of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of sequences in the control sample as compared with the relative copy numbers of sequences in the carray in determined using an intercalating does or a detectable antibody, or its frammer that is geneific for a mucleic acid detectable antibody or is detectable.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable moiety such that hybridization of the genomic array is determined by determining the presence, absence, amount or location of the detectable label on the one or more genomic arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence represents a non-nucleotide probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kirtsen NV, Hyldig-Nielsen JJ, Williams BF;
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DAKOCYTOMATION DENMARK AS.
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Gaps

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0; Indels

Length 20;

Score 20; DB 1; Le Pred. No. 6.8e+02;

100.0%; Pre-

Query Match Best Local Similarity

Conservative

Mismatches

2709 AAAAAAAAAAAAAAAA 2728

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1 AAAAAAAAAAAAAAA 20

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The present sequence represents a non-nucleotide probe. The probe is useful for suppressing the binding of one or more detectable nucleic acid probes, that are greater than 100 base pairs and that have been derived from genomic nucleic acid, to one or more undesired sequences in an assay for determining the sample with the mixture of probes in preferably comprises contacting the sample with the one or more detectable nucleic acid probes, and determining the target genomic nucleic acid probes, on determining the hybridization of the one or more detectable nucleic acid probes to the target genomic nucleic acid of the sample by determining the hybridization of the one or nucleic acid of the sample is metaphase spreads, interphase nucleic acid of the sample is metaphase spreads, interphase nucleic acid of the sample is metaphase spreads, interphase nucleic acid with the tart of a control sample using a sample of genomic nucleic acid with the the one or nucleic acid, which are differentially labelled, the array with the mixture of the probe under suitable hybridization conditions, and comparing the array with treated mixture conditions, and comparing the array with treated mixture of sample and control genomic nucleic acid under suitable hybridization conditions, and comparing the intensities of the signals from the differential labels of the array to that caused by hybridization of the probes to genomic nucleic acid, thus determining one or more variations in copy numbers of substantially each acid, thus sample as compared with the relative copy numbers of substantially
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Non-nucleotide probe for suppressing binding of detectable nucleic acid probes to undesired sequences, has aggregate nucleobase sequence homologous to randomly distributed repeat sequence of genomic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                        non-nucleotide probe; Bacterial Artificial Chromosome clone; BAC; 88;
                                                                            Gaps
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                                  DB 1; Length 20;
6.8e+02;
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Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                    0.7%; Score 20; DB
100.0%; Pred. No. 6.E
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                    Non-nucleotide probe of the invention #24
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DAKOCYTOMATION DENMARK AS.
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                    Query Match
Best Local Similarity 100.v
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
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ö Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiaethmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or identical sequences in the control. The hybridization of the genomic array is determined using an intercalating dye or a detectable antibody, or its fragment, that is specific for a nucleic acid/nucleic acid hybrid. The sample of genomic nucleic acid to be tested and the reference of nucleic acid are labelled with detectable moiety such that hybridization of the genomic array is determined by determining the presence, absence, amount or location of the detectable label on the one or more genomic arrays. The genomic array comprises nucleic acid that is prepared from Bacterial Artificial Chromosome (BAC) clones. The present sequence represents a non-nucleotide probe of the invention. Human, antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatesthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine esensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. **Gaps** Aguilar D; ö Length 20; 0; Indels Katz E, Pabalan J, Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 0.7%; Score 20; DB 1; Lv 100.0%; Pred. No. 6.8e+02; ative 0; Mismatches 0; Disclosure; SEQ ID NO 3509; 872pp; English. ŝ 1 AAAAAAAAAAAAAAAAAA 20 Li Y, Sandrasagra A, Tang L, Shahabuddin Human oligonucleotide sequence. BP. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC. ABZ88267 standard; DNA; 20 (first entry) 20; Conservative WPI; 2003-229219/22. Best Local Similarity WO200285308-A2. Homo sapiens. 17-OCT-2003 31-OCT-2002. Nyce JW, ] Miller S, ubiquinone ABZ88267; Query Match RESULT 492 ABZ88267 Matches 8888888888888 8

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchocometriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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Gaps
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  Length 20;
                         0; Indels
 0.7%; Score 20; DB 1; Le
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0;
                                                   2709 AAAAAAAAAAAAAAAA 2728
                                                                            1 AAAAAAAAAAAAAAA 20
                         20; Conservative
             Local Similarity
Query Match
                         Matches
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## RESULT 493

Human oligonucleotide sequence. ABZ88565 standard; DNA; 20 BP (first entry) 17-OCT-2003 ABZ88565; 

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

ä Pabalan J, Aguilar щ Katz Sandrasagra A, Ka i, Shahabuddin S; WPI; 2003-229219/22. Li Y, Sar Tang L, Miller S, Nyce JW,

Pharmaceutical composition for treating ailments associated with impaired

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or ö The invention relates to a novel pharmaceutical composition, which has respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid Disclosure; SEQ ID NO 3807; 872pp; English. ubiquinone

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this parent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
                         0; Indels
                         0; Mismatches
            100.08;
                         20; Conservative
            Similarity
  Query Match
               Local
                         Matches
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## 2709 AAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAA 20

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Human oligonucleotide sequence. BP. ABZ88619 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ88619; 494 RESULT 49 ABZ88619 

Human; antisense; lung dysfunction; nasal airway dysfunction; antializammatory; sucroid; ubiquinone; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

ä Pabalan J, Aguilar Katz E, Sandrasagra A, Ka , Shahabuddin S; Li Y, San , Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 3861; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, cintialsthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from MIPO at ftp.wipo.int/pub/published_pct_sequences
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Gaps ; 0 0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. Mc-1144 0; Mismatches Conservative Query Match Best Local Similarity Matches 20; Conserv ઠે

2709 AAAAAAAAAAAAAAAA 2728 

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RESULT 495

ABZ89705 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ89705; 

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; phypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

sapiens Ношо

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; WPI; 2003-229219/22. Nyce JW, I Miller S,

Pharmaceutical composition for treating ailments associated with impaired

respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid

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ubiquinone.

Disclosure; SEQ ID NO 4947; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genee encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an anticinflammatory steroid in a subject, for reducting or depleting levels of, or reducing sensitivity to adenosine, reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Gaps ö Length 20; 0; Indels Query Match 0.7%; Score 20; DB 1; L4 Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0;

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RESULT 496

ABZ88816 standard; DNA; 20

ВР.

ABZ88816; 

(first entry) 17-0CT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Aguilar D; Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4058; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiashmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory eterot or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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   Length 20;
0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                          2709 AAAAAAAAAAAAAAA 2728
                                                                                       1 AAAAAAAAAAAAAAA 20
                             20; Conservative
            Local Similarity
Query Match
                             Matches
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RESULT 497 ABZ88881

ABZ88881 standard; DNA; 20 ABZ8881: 

BP.

(first entry) 17-OCT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-0CT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar Katz E, Tang L, Shahabuddin S; Sandrasagra A, Li Y, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired or respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Disclosure; SEQ ID NO 4123; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or The invention relates to a novel pharmaceutical composition, which has

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPPO ftp.wipo.int/pub/published\_pct\_sequences

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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. ... Matches 20; Conservative Similarity Query Match Local

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Human oligonucleotide sequence. ABZ89706 standard; DNA; 20 BP (first entry) 17-0CT-2003 ABZ89706; RESULT 498 **ABZ89706** 

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

ä Pabalan J, Aguilar щ Katz Sandrasagra A, Li Y, San Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Disclosure; SEQ ID NO 4948; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fip.wipo.int/pub/published_pct_sequences
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Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
                      0; Indels
   100.0%; Pred. No.
  7%; Score 20;
                         Conservative
  Query Match
Best Local Similarity
Matches 20; Conserv
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2709 AAAAAAAAAAAAAAAAA 2728 

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Human oligonucleotide sequence.
   ABZ88620 standard; DNA; 20 BP
            17-OCT-2003 (first entry)
       ABZ88620;
RESULT 499
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Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatesthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchodonstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Sandrasagra A, Ka L, Shahabuddin S; Li Y, Sar Tang L, Nyce JW, L Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 3862; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiniflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallerial antiachmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiniflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Gaps ö Length 20; 0; Indels Ouery Match 0.7%; Score 20; DB 1; Lv Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0;

## 2709 AAAAAAAAAAAAAAAAA 2728 20

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BP. ABZ88880 standard; DNA; 20 17-OCT-2003 (first entry) ABZ88880; RESULT 500 ABZ88880 

Human oligonucleotide sequence.

Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

ä Aguilar Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4122; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiaethmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung suffectent in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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ABZ89179 standard; DNA; 20 BP. (first entry) 17-OCT-2003 ABZ89179; RESULT 501 ABZ89179 

ABZ88814 standard; DNA; 20 BP

502

ABZ88814

(first entry)

17-OCT-2003

ABZ88814;

Human oligonucleotide sequence.

Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar Katz E, Sandrasagra A, Ka i, Shahabuddin S; Li Y, Sar Tang L, Miller S, Ŗ,

ä

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(8) antisense to specific gene(8) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4421; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostetic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or invention relates to a novel pharmaceutical composition, which has

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this parent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                 Sequence 20 BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                    100.0%; Pred. No.
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Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiantiasthmatic; hypotensive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human oligonucleotide sequence. 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. WO200285308-A2 Homo sapiens 31-OCT-2002. 

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; WPI; 2003-229219/22. Miller S, Nyce JW,

(EPIG-) EPIGENESIS PHARM INC

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4056; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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7%; Score 20; DB 1; Length 20;
                                       0; Indels
                      6.8e+02;
  Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Best Local Solicative 0; Mismatches 20; Conservative 0; Mismatches
                                                                                             2709 AAAAAAAAAAAAAAAAAA 2728
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Human oligonucleotide sequence.
   ABZ89241 standard; DNA; 20
           (first entry)
           17-OCT-2003
       ABZ89241;
RESULT 503
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiliflammatory steroid; ubiquinone; antiliflammatory; antiallergic; antiaefhmatic; Mypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

WO200285308-A2. sapiens Ношо

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Nyce JW, L Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4483; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinifiammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung allengies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                   88666666666888888
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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                      Gaps
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   Length 20;
                      0; Indels
   Score 20; DB 1; Le
Pred. No. 6.8e+02;
         100.0%; Pred. ....
    0.7%;
Ouery Match
Best Local Similarity 100.8
Matches 20, Conservative
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Gaps

ABZ90650 standard; DNA; 20 (first entry) 17-OCT-2003 AB290650; RESULT 504 ABZ90650 

ВР

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

ä Katz E, Pabalan J, Aguilar ŝ Sandrasagra A, L, Shahabuddin Li Y, Sand, Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone

Disclosure; SEQ ID NO 5892; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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8888888888888
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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed profice of the printed printed profice of the printed printed profice of the printed profice of the printed profice of the printed profice of the printed printed profice of the printed profice of the profice of the printed profice of the profice of the printed profice of the printed profice of the printed profice of the printed profice of the at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Length 20; 0; Indels Score 20; DB 1; Le Pred. No. 6.8e+02; 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAA 20 0.7%; 20; Conservative Local Similarity Query Match Matches ð a

ABZ88815 standard; DNA; 20 BP RESULT 505 ABZ88815

(first entry) 17-OCT-2003 ABZ88815; 

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

24-APR-2001; 2001US-0286137P

23-APR-2002; 2002WO-US013135.

(EPIG-) EPIGENESIS PHARM INC.

ä Pabalan J, Aguilar Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired or respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Disclosure; SEQ ID NO 4057; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or The invention relates to a novel pharmaceutical composition, which has

for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPPO preventing a respiratory, lung or malignant disease or condition, also ftp.wipo.int/pub/published\_pct\_sequences

88888888888888

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; .00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Prec. Matches 20; Conservative Similarity Query Match Best Local

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Gaps

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ABZ85311 standard; DNA; 20 BP RESULT 506 ABZ85311

(first entry) 17-OCT-2003

ABZ85311;

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; cimmunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

ä Katz E, Pabalan J, Aguilar Sandrasagra A, Ka , Shahabuddin S; Li Y, Sar Tang L, Miller S, Nyce JW, 

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 553; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an attinfilammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 8X8888888888X8

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

Gaps ö Query Match 0.7%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAA 1 셤 ઠે

Human oligonucleotide sequence. BP. ABZ85435 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ85435; RESULT 507 

antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antiaense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. antisense; lung dysfunction; nasal airway dysfunction; Human;

Homo варіепв.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Li Y,

ä

Aguilar

Pabalan J,

Sandrasagra A, Katz E, , Shahabuddin S;

Tang L,

Nyce JW, 1 Miller S,

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or WPI; 2003-229219/22.

Claim 15; SEQ ID NO 677; 872pp; English.

ubiquinone.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genee encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

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Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

ö Gaps ö Length 20; 0; Indels 0.7%; Score 20; DB 1; L. 100.0%; Pred. No. 6.8e+02; Live 0; Mismatches 0; Local Similarity 100. nes 20; Conservative Query Match Matches

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RESULT 508 ABZ88817

BP.

ABZ88817 standard; DNA; 20 ABZ88817; 

Human oligonucleotide sequence.

17-OCT-2003 (first entry)

Human, antisense; lung dysfunction, nasal airway dysfunction; antisinflammatory steroid; ubiquinone; antisinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Katz E, Pabalan J, ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Li Y, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4059; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory, steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory effect of an antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of uniquininone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Gaps
                      ;
  Length 20;
                     Indels
Score 20; DB 1; Le
Pred. No. 6.8e+02;
     100.0%; Preα....
ive 0; Mismatches
                                         2709 AAAAAAAAAAAAAAA 2728
                                                             1 AAAAAAAAAAAAAAA 20
 0.7%;
        Local Similario,
nes 20; Conservative
 Query Match
                    Matches
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ABZ88939 standard; DNA; 20 BP (first entry) 17-OCT-2003 ABZ88939; RESULT 509 ABZ88939 

Human oligonucleotide sequence

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Sandrasagra A, Ka i, Shahabuddin S; WPI; 2003-229219/22. Tang L, Li Y, ŝ Nyce JW, Miller

Pharmaceutical composition for treating ailments associated with impaired

respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid

or

Disclosure; SEQ ID NO 4181; 872pp; English.

ubiquinone.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or invention relates to a novel pharmaceutical composition, which has a

preventing a respiratory, lung or malignant disease or condition, also antin the prophylactic or therapeutic respiratory effect of an antinfilammatory steroid in a subject, for reducing or despleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchocomstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this parent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences 

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; .00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Prec. ... 20; Conservative Local Similarity Query Match Matches

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2709 AAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAA 20 ઠે 요

멾 ABZ89302 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ89302; 210 RESULT 51 ABZ89302 

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory; antiallergic; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

ä Katz E, Pabalan J, Aguilar ŝ Sandrasagra A, Li Y, Sar Tang L, Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(8) antisense to specific gene(8) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4544; 872pp; English.

first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or The invention relates to a novel pharmaceutical composition, which has a

for enhancing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing or adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences 888888888888888888

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches 2709 AAAAAAAAAAAAAAA 2728 8

1 AAAAAAAAAAAAAAA 20 g

ABZ88566 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ88566; RESULT 511 

BP.

Human oligonucleotide sequence.

antinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; antisense; lung dysfunction; nasal airway dysfunction; inflammation; respiratory disease; ds. Human; lung

Homo sapiens

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, S; Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 3808; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antichilammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published\_pct\_sequences

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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Gaps ö Length 20; 0; Indels 0.7%; Score 20; DB 1; L 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Best Local Similarity 100. Matches 20; Conservative Query Match

2709 AAAAAAAAAAAAAAAAA 2728 

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Gaps

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512 ABZ89086 RESULT

ABZ89086 standard; DNA; 20

BP.

ABZ89086; 

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

Human, antisense, lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone, antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

Aguilar D;

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4328; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genemic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing or depleting levels of, or reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Length 20; 0; Indels Score 20; DB 1; Le Pred. No. 6.8e+02; 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAA 2728 0.7%; 20; Conservative Similarity Query Match Local Matches 8 셤

1 AAAAAAAAAAAAAAA 20

ABZ85533 standard; DNA; 20 BP 17-OCT-2003 ABZ85533; RESULT 513 

(first entry)

Human oligonucleotide sequence,

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunouppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds

Homo sapiens

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Katz E, Sandrasagra A, Li Y, Nyce JW,

Tang L, Shahabuddin S;

Miller S,

ä

Pabalan J, Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired or respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Claim 15; SEQ ID NO 775; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or invention relates to a novel pharmaceutical composition, which has a

preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this parent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Score 20; DB 1; Length 20; 0; Indels 100.0%; Pred. No. 6.8e+02. 0.7%; 20; Conservative Similarity Query Match Local Matches

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2709 AAAAAAAAAAAAAAAA 2728 

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Gaps

ABZ89015 standard; DNA; 20 (first entry) 17-0CT-2003 ABZ89015; 514 ABZ8901 

BP

Human; antisense; lung dysfunction; nasal airway dysfunction; antialflammatory steroid; ubiquinone; antianflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human oligonucleotide sequence.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

ä Katz E, Pabalan J, Aguilar Sandrasagra A, Ke L, Shahabuddin S; Li Y, Sar Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4257; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 8888888888888888

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

.7%; Score 20; DB 1; Length 20; 3.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAAAA 20 Query Match 0.7 Best Local Similarity 100. Matches 20; Conservative 8

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Human oligonucleotide sequence. ABZ89240 standard; DNA; 20 (first entry) 17-OCT-2003 ABZ89240; RESULT 515 

BP

Human, antisense; lung dysfunction, nasal airway dysfunction, antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Ношо варіепя

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

ä Pabalan J, Aguilar Katz E, S; i Y, Sandrasagra A, Tang L, Shahabuddin Li Y, Nyce JW, L Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure, SEQ ID NO 4482; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or The invention relates to a novel pharmaceutical composition, which has a

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

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Sequence 20 BP; 18 A; 1 C; 0 G; 1 T; 0 U; 0 Other;

ö Gaps ; Length 20; 0; Indels 0.7%; Score 20; DB 1; L 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; 20; Conservative Best Local Similarity Query Match Matches

2707 CTAAAAAAAAAAAAAAA 2726 

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Gaps

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ABZ89441 standard; DNA; 20 RESULT 516 ABZ8944] 

ВЪ.

ABZ89441;

17-OCT-2003 (first entry)

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Human oligonucleotide sequence.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Katz E, Pabalan J, ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Li Y, Miller S, Nyce JW,

ä

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4683; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiachmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, lunt was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Length 20; 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels 2709 AAAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAA 20 Local Similario, nes 20; Conservative Query Match Matches ò 요

RESULT 517

ABZ89016 standard; DNA; 20 BP

ABZ89016;

(first entry) 17-0CT-2003 Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antianflammatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; inflammation; respiratory disease; ds. lung

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its s) antisense to specific gene(s) or its glucocorticoid or non-glucocorticoid steroid respiration, has oligo(s) corresponding RNAs, and gl ubiquinone.

Disclosure; SEQ ID NO 4258; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genee encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostetic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or invention relates to a novel pharmaceutical composition, which has a 

preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences 88888888888888888

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 0; Indels Pred. No. 6.8e+02; 100.0%; Pred. .v.. 20; Conservative Similarity Query Match Local Matches

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2709 AAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAAA

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Gaps

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518 ABZ89120

ABZ89120 standard; DNA; 20

ABZ89120:

BP

(first entry) 17-OCT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antianflammatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Katz E, Pabalan J, Aguilar Sandrasagra A, Ka L, Shahabuddin S; Li Y, San Tang L, Miller S, Nyce JW. 

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WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4362; 872pp; English.

first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end ganomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or The invention relates to a novel pharmaceutical composition, which has a

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preventing a respiratory, lung or malignant disease or condition, also active enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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Score 20; DB 1; Length 20; Pred. No. 6.8e+02;
                        0; Indels
     100.0%; Pred. ...
  7%; Score 20;
                                                   2709 AAAAAAAAAAAAAAAAA 2728
                           Conservative
  Query Match
Best Local Similarity
Matches 20; Conserv
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Gaps

1 AAAAAAAAAAAAAAAA 20 셤

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Human oligonucleotide seguence.
   ABZ89704 standard; DNA; 20
            (first entry)
            17-OCT-2003
        ABZ89704;
RESULT 519
 AB289704
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Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatethmatic; propertait; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

WO200285308-A2. sapiens 31-OCT-2002. Homo

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, s; Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, I Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4946; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or

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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antichilammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition, love: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nanotechnology, ss, nucleic acid detection, nanoparticle; virus detection; human immunodeficiency virus; HIV; hepatitis; herpes; cytomegalovirus; Epetein-Barr virus; bacterial disease; DNA sequencing; sexually transmitted disease; inherited disorder; forensic; paternity testing; cell line authentication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting nucleic acids having 2 portions e.g. for detecting disease, comprises use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the nucleic acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel method for detecting a nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nanotechnology nucleic acid detection method associated #54.
                                                                                                                                                                                                        Length 20;
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                                                                                                                                                                          Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                      Score 20; DB 1; I
Pred. No. 6.8e+02;
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/note= "OTHER= Thiol modified"
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                                                                                                                                                                                                                                                                                       BP.
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29-JAN-1999; 9US-00240755.
25-JUN-1999; 9US-00344667.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
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                                                                                                                                                                                                                                                                                                                                                                             ACD27320 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-228115/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                            ACD27320;
                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Taton
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having 2 portions. The method comprises providing nanoparticles having oligonucleotides attached, where the oligonucleotide on each nanoparticle has a sequence of 2 portions of nucleic acid.

The nucleic acid and nanoparticle are contacted to allow hybridisation of the oligonucleotide on the nanoparticle with two or more portions of nucleic acid and observing a detectable change brought about by the chybridisation. The method of the invention is useful for separating a selected nucleic acid having 2 portions. The method of the invention is useful for detecting nucleic acids which may be convented in sequencing of nucleic acids which may be referrably, the method of disease and in sequencing of nucleic acids which may be preferrably, the method is useful for detecting nucleic acids for diseases and in sequencing of nucleic acids for cidagnosis and/or monitoring of viral diseases (human immunodeficiency virus, bepatitis virus, herpes virus, cytomegalovirus and Epstein-Barr covirus, bacterial diseases, sexually transmitted diseases, inherited diseases, in DNA sequencing, for paternity testing, for call line authentication, for monitoring gene therapy, etc. This method cidagnosis and one passive detecting nucleic acids based on observing a colour change with the naked eye so is cheap, fast, simple and robust, and does not require secientised expensive equipment. The present sequence represents a thiol increase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Probe; nucleic acid detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC58867 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-SEP-2003' (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Doubly labelled DNA probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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The present invention describes a method (M1) for preparing an internucleotide phosphorothioate linkage enriched in the Sp or Rp anationer between a synthon having a hydroxyl moiety at the 5' position and a 2'-substituted nucleoside having an activated phosphate moiety at the 3'-position, comprising coupling a synthon with a 2'-substituted nucleoside in the presence of coupling agent that is selected to enhance either the Rp or Sp enantiomer according to its pKa. This method is useful for preparing an oligonucleotide having at least one region of internucleotide linkages that is enhanced in the Sp or Rp enantiomer, which involves providing a nucleotide chain having a hydroxyl moiety at the 5'-position or a growing oligonucleotide chain having a hydroxyl moiety at the 5'-position, coupling the nucleotide or growing oligonucleotide chain to a 2'-substituted nucleoside having an activated phosphate moiety at
                                                                                                                                                                                                                                 Preparing internucleotide phosphorothioate linkage enhanced in Sp/Rp enantiomer, by coupling a synthon with 2' substituted nucleoside in presence of coupling agent having a pKa that enhances linkage in Sp/Rp enantiomer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 31; 65pp; English.
                                                                                                             WPI; 2003-157021/15.
Ravikumar VT;
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The present sequence is an example of nucleic acid probes of the invention. The probe may be doubly labelled with non-identical covalently attached dyes, e.g. the fluorescent intercalator tehidium, which serves as the detector dye and the fluorescent dye fluorescein, which serves as the donor dye of a fluorescent resonance energy transfer (FRET) system. A

Detection or quantification of nucleic acid analyte, by hybridizing a nucleic acid probe having non-identical covalently attached dyes, with nucleic acid analyte, and measuring change in fluorescence of the probes.

Wolter A;

Davies M,

Bruce I,

(PROL-) PROLIGO LLC.

WPI; 2003-505122/47.

Example 9; Page 32; 110pp; English.

ö bifunctional linker was used to attach the dyes to the oligonucleotide. The probe generates a fluorescent signal upon hybridisation to a complementary nucleic acid based on the interaction of the intercalator with the formed double-stranded DNA. Nucleic acid probes of the invention can be used in homogeneous assays, real-time PCR monitoring, transcription assays, expression analysis on nucleic acid microarrays and other microarray applications such as genotyping Gaps Chiral; phosphorothioate; oligonucleotide synthesis; enantiomer; ss. ö 0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels 1. .20 /\*tag= a /mod\_base= OTHER /note= "phosphorothioate linkages" Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; 100.0%; Prec. ... Phosphorothioate 20-mer oligonucleotide #1. Location/Qualifiers 2709 AAAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAA BP. 13-JUN-2002; 2002WO-US018581. 14-JUN-2001; 2001US-00881535. ABZ22916 standard; DNA; 20 (first entry) 20; Conservative (ISIS-) ISIS PHARM INC. Similarity WO2002102815-A2 Key modified\_base 08-APR-2003 27-DEC-2002. Synthetic. ABZ22916; Query Match Local Matches ABZ22916, 888888888888 ò g 

ö the 3' position in the presence of the coupling agent, and repeating the coupling step until the desired number of linkages is established. The oligonucleotide having a region of internucleotide linkages that is enhanced in the Sp enantiomer is further processed to include another region of internucleotide linkages that is enhanced in the Sp and/or Rp enantiomer. Oligonucleotides prepared by the method lead to improved drugs, diagnostics and research reagents. The present sequence represents an oligonucleotide used in the exemplification of the present invention Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distance; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Gaps Pharmaceutical composition for treating asthma, has antisense of algonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. Pabalan J, Aguilar D; o; 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; pulmonary transplantation rejection; ss; primer. AI652901-derived oligonucleotide SEQ ID 3509. ы Li Y, Sandrasagra A, Katz Tang L, Shahabuddin S; 2709 AAAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAAA ВЪ 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC. ABD24497 standard; DNA; 20 (first entry) Conservative WPI; 2003-093058/08. Local Similarity nes 20; Conserv WO200285309-A2. Homo sapiens 31-OCT-2002. 29-JUL-2004 Nyce JW, I Miller S, ABD24497; 20 Query Match RESULT 523 Matches ABD24497 8888888888 δ g

instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-darenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered correction are associated composition comprises oligo and is administered to reduce the production or reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung correction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, cinflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it ö Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Gape Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and Aguilar D; ö Length 20; IndelB E, Pabalan J, Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; ö 0.7%; Score 20; DB 1; L. 00.0%; Pred. No. 6.8e+02; AI128305-derived oligonucleotide SEQ ID 4059. 100.0%; Pred. .... Katz S; 2709 AAAAAAAAAAAAAAAAA 2728 Li Y, Sandrasagra A, Tang L, Shahabuddin BP. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC. ABD25047 standard; DNA; 20 (first entry) Conservative bronchodilating agent WPI; 2003-093058/08. Best Local Similarity WO200285309-A2. Homo sapiens. 29-JUL-2004 31-OCT-2002. 20; Nyce JW, I Miller S, ABD25047; Query Match RESULT 524 Matches ABD25047 셤 ଟ

comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c)

This invention describes a novel composition (a) a first active agent,

Claim 15; SEQ ID NO 3509; 763pp; English.

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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analygesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung confinemation, allergies, asthma, impeded respiration, respiratory disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, confinemation, emphysema, chronic obstructive pulmonary disease, pulmonary treaduced adenosine content of the anti-sense oligos corresponding to the mynicianes present in the target RNA serves to prevent the breakdown of the pulmonary the opinonic operation or prevent the breakdown of the pulmonary the opinonic operation or prevent the present in the target RNA serves to prevent the breakdown of the pulmonary the opinonic portent of the encodence of prevents in the target RNA serves to prevent the present in the target RNA serves to prevent th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby,
Claim 15; SEQ ID NO 4059; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prevent any unwanted effects due to it
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels 0.7%; 100.0%; Pre-Conservative Similarity 20; Query Match Local Matches

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ABD25316 standard; DNA; 20 BP (first entry) 29-JUL-2004 ABD25316; RESULT 525 ABD2531 

AI092429-derived oligonucleotide SEQ ID 4328.

beta-adrenergic agonist, respiratory disease; pulmonary vasoconstriction, respiratory distress syndrome, allergic rhinitis; pulmonary hypertension, emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity, lung, cancer, surfactant depletion, antiallergic, antiinflammatory; antiasthmatic, analgesic, hypotensive, immunosuppressive, cytostatic, cystic fibrosis, pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, ıdrasagra A, K. Shahabuddin S; Sandrasagra Li Y, San Tang L, Miller S, Nyce JW,

WPI; 2003-093058/08

ဌ oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing bronchodilating agent.

Claim 15; SEQ ID NO 4328; 763pp; English.

This invention describes a novel composition of a first active agent, comprising oligomucleotides, effective for alleviating brunchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and corresponding adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligomucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) device, in separate containers, (b) the oligomucleotides, (c) analgesic, hypotensive, immunosuppressive and cytostatic activity, is a containers agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cyllmonary obstruction, and/or bronchoconstriction and/or lung with a disease or conditions and/or bronchoconstriction and associated with a disease or conditions and by a subman, impeded respiration, respiratory cillammation, allergies and/or bronchoconstriction and associated inflammation, allergies, asthma, impeded respiration, respiratory contemporation rejection, pulmonary bulmonary disease, pulmonary contemplantation rejection, pulmonary bereves to prevent the breakdown of the infinite present in the target RNA serves to pervent the breakdown of the change or conditions are separated by a prevent or pervent to be a process of the conditions are associated or the conditions the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it invention describes a novel composition (a) a first active agent,

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. ... 20, Conservative Similarity Query Match Best Local S Matches

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Gaps

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ABD21763 standard; DNA; 20 BP. (first entry) 29-JUL-2004 ABD21763; RESULT 526 ABD21763 

Human stanniocalcin-derived oligo SEQ ID 775.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;

antisense; bronchoconstriction; allergy; hyposecretion; pain;

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ABD25246 standard; DNA; 20 BP.
                                                         ABD25246;
RESULT 527
                ABD25246
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respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and AI051839-derived oligonucleotide SEQ ID 4258. Claim 15; SEQ ID NO 4258; 763pp; English. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC. 29-JUL-2004 (first entry) bronchodilating agent. WPI; 2003-093058/08 WO200285309-A2. Homo sapiens. 31-OCT-2002. Miller S, Nyce JW, Human: This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchconstruction, respiratory tract inflammation, alleviating bronchconstruction, respiratory tract inflammation, alleviates and censitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention as antiallergic, antiafflammatory, antiathmatic, an angesic, hypotensive, immunosuppressive and cytostatic activity, is beta-adrenergic agonist. The composition is useful for preventing or reasting a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted tnucleic acids associated with lung airway or lung dysfunction, and Katz E, Pabalan J, Aguilar D; Score 20; DB 1; Length 20; Pred. No. 6.8e+02; Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; pulmonary transplantation rejection; ss; primer Claim 15; SEQ ID NO 775; 763pp; English. Sandrasagra A, Ka ,, Shahabuddin S; 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC 0.7%: Query Match 0.7 Best Local Similarity 100. Matches 20; Conservative bronchodilating agent. Li Y, Sar Tang L, WPI; 2003-093058/08 WO200285309-A2. Homo sapiens 31-OCT-2002 ŝ Nyce JW, Miller

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Aguilar D;

Katz E, Pabalan J, S;

Sandrasagra A,

Li Y, Sand Tang L, (

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating cadenosine sensitivity, levels of adenosine (A) or (A) receptors, reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung arway or lung expression or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) confociations for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, is a conficient or malgesic, hypotensive, immunosuppressive and cytostatic activity, is a confociation are application comprises oligo and is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The culmonary obstruction, and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary vasconstriction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, distress syndrome, pain, cystic fibrosis, allergic thinits, pulmonary cor transplantation rejection, pulmonary infections, brundings present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the present of the oligonucleotides into products that free adenosine nor the oligonucleotides into products that free adenosine or the oligonucleotides into products that free adenosine or the oligonuc kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it heart, e.g.,

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Gaps

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Mismatches

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DB 1; Length 20; Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; Score 20; Query Match Best Local Similarity

100.0%; Pred. No. 6.8e+02;

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 Gaps
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Conservative
20;
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2709 AAAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAA 20 셤

ABD24849

BP ABD24849 standard; DNA; 20

ABD24849;

(first entry) 29-JUL-2004

AI092623-derived oligonucleotide SEQ ID 3861.

analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;

Homo sapiens.

WO200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

ä Aguilar Pabalan J, Katz E, Sandrasagra A, Ka L, Shahabuddin S; Tang Ľi Y, Miller S, 3 Nyce

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 3861; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating by the comprising oligonucleotides, effective for alleviating construction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cuffactant depletion or hyposecretion, when administered to a marmal. The oligonucleotides are derived from a gene encoding or regulating corresponding or regulating expression of a target polypeptide associated with lung airway or lung dystunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a containers, instruction is useful for preventing or treating a respiratory, lung or malignant disease. The administered conventing or treating a respiratory, lung or malignant disease. The administered conventing or treating a respiratory, lung or malignant disease. The administered convention, and/or bronchoconstriction and/or lung confiction and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergies plantatory. 

ö the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of Gaps ö Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 0.7%; Sco... 100.0%; Pred. No. v... 0; Mismatches 2709 AAAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAA 20; Conservative Similarity Query Match Local Matches 888888888 ò g

AI041212-derived oligonucleotide SEQ ID 4482. ВР ABD25470 standard; DNA; 20 (first entry) 29-JUL-2004 ABD25470; RESULT 529 ABD25470 

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; lumunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory districtses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

sapiens. Ношо WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar Pabalan J, Katz E, Sandrasagra A, Ka i, Shahabuddin S; Li Y, Sar Tang L, Miller S, Nyce JW,

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WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4482; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or resplating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides (c) instructions for adding a carrier and for use of the kit. The composition

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analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mENA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung or inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated vith a disease or condition such as pulmonary vasoconstriction, or inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary cransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonuclectides into products that free adenosine into the system of prevent any unwanted effects due to it
invention has antiallergic, antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 18 A; 1 C; 0 G; 1 T; 0 U; 0 Other;
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DB 1; Length 20; 6.8e+02; 0; Indels 100.0%; Pred. No. 6.8 :ive 0; Mismatches 0.7%; Score 20; Local Similarity 100. nes 20; Conservative Query Match Matches

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Gaps

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2707 CTAAAAAAAAAAAAAA 2726 CTAAAAAAAAAAAAAA 20 ઠે 엄

ABD21665 standard; DNA; 20 BP 29-JUL-2004 (first entry) ABD21665; RESULT

Human stanniocalcin-derived oligo SEQ ID 677.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

W0200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, Sandrasagra A, Ka , Shahabuddin S; Li Y, San Tang L, Nyce JW, I Miller S,

WPI; 2003-093058/08.

ದಿ Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted truncleic acids associated with lung airway or lung dysfunction, and oronchodilating agent.

Claim 15; SEQ ID NO 677; 763pp; English

This invention describes a novel composition, alleviating describes and describes and comprishing oligomucleotides, effective for alleviating comprishing adenosine sensitivity, levels of adenosine (A) or (A) receptors, reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligomucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction also describes a kit, that comprises: (a) a delivery of the invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallersic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytosetaides, (c) that invention has antiallersic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytosetaid or treating or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactant hypoproduction and/or lung corresponding such a disease or condition such as pulmonary vasoconstriction, composition rejection, pulmonary infections, bronchises uplanonary disease, pulmonary cyclopertension, emphysema, chronic obstructive pulmonary energet maneral such cyclopertension and cyclopertension and cyclope This invention describes a novel composition (a) a first active agent, brain, heart, kidney, etc, unwanted effects due to it prevent any X38888888888888888888888888888888888

ö Length 20; 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; DB 1; Le 0.7%; Score 20; DB 100.0%; Pred. No. 6.8 ive 0; Mismatches Local Similarity 100. Les 20, Conservative Query Match Matches

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2709 AAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAA 1 ð 셤

ABD24796 standard; DNA; 20 BP. ABD24796; RESULT 531 ABD24796 

29-JUL-2004 (first entry)

All22689-derived oligonucleotide SEQ ID 3808.

Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D;

Katz E,

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating compression of a target polypeptide associated with lung airway or lung thvention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, and adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, and adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition, comprises oligo and is administered to reduce the production or variability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, allergies and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary vasconstriction, inflammation, allergies, asthma, impeded respiratory disease, pulmonary obstruction, mephysema, chronic obstructive pulmonary disease, pulmonary cranger condition, pulmonary incedections, benchmany incedections, benchmany incedections, benchmany incedection, pulmonary incedections, benchmany incedections, pulmonary incedections, benchmany in the target RNA serves to prevent the breakdown of the oligonucleotides into product that free adenosine into the system of the oligonucleotides into product that the term and unawanted effects due to it transplante
                                                                                                                                                                                          oligonucleotide containing less percentage of adenosine, targeted t
nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 3808; 763pp; English.
                            ii Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                      bronchodilating agent.
                                                                                                         WPI; 2003-093058/08
                            Li Y,
                                                            Miller S,
                            Nyce JW,
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö , DB 1; Lens. 7, 6.8e+02; 0; Indels 0.7%; Score ... 100.0%; Pred. No. 6.8e-Query Match
Best Local Similarity 100.

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ઠ g RESULT 532 ABD25045

ABD25045 standard; DNA; 20 BP. 

ABD25045;

AI128305-derived oligonucleotide SEQ ID 4057.

29-JUL-2004 (first entry)

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthantic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rihinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Aguilar D; Pabalan J, Katz E, Li Y, Sandrasagra A, K Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC WPI; 2003-093058/08. WO200285309-A2. Homo sapiens. 31-OCT-2002 Miller S, 

ဌ Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted truncleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4057; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configurated from a gene encoding or regulating or expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises; (a) a delivery cancering for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition configuration. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition configuration and or and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary observation, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, cancer. Inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary disease, pulmonary disease, pulmonary of transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense observent the breakdown of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine environment and thereby, to intervant any unwanted effects due to prevent any unwanted effects due to it

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; Indels ö 100.0%; Pred. n. 20; Conservative Local Similarity Query Match Matches

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2709 AAAAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAA 20 BP. RESULT 533 ABD25350 셤 8

ABD25350 standard; DNA; 20 SXXX

ABD25350

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(EPIG-) EPIGENESIS PHARM INC 

Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. AI096522-derived oligonucleotide SEQ ID 4362. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (first entry) WO200285309-A2. Ното варіепв 31-OCT-2002 29-JUL-2004

oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. Katz E, Pabalan J, Aguilar D; Pharmaceutical composition for treating asthma, has antisense Sandraвagra A, Ka , Shahabuddin S; WPI; 2003-093058/08. Tang L, Li Y, Nyce JW, L Miller S,

Claim 15; SEQ ID NO 4362; 763pp; English

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The cligonucleotides are derived from a gene encoding or regulating cyptemotion or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or mailganat disease. The administered composition comprises oligo and is administered to reduce the production or raduce the amount of target polypeptide present in the lungs. The culting mandion, allergies and/or bronchoconstriction and a sesociated inflammation, allergies and/or bronchoconstriction are associated inflammation, allergies and/or surfactant hypopropurching are associated inflammation, allergies and/or surfactant hypopropurching are associated inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypetrension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thy discounties present in the target RNA serves to prevent the breakdown of the oligomuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it with a disease or condition such as pulmonary vasoconstriction,

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

O.7%; Score 20; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 6.8e+02; Local Sol, Conservative 0; Mismatches 0; Indels Query Match Matches

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Gaps

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AAAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAA 20 2709 ò 원

RESULT 534

ABD25245 standard; DNA; 20 ABD25245

ABD25245;

29-JUL-2004 (first entry)

AI051839-derived oligonucleotide SEQ ID 4257.

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; oystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vosoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

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24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Miller S, 3

WPI; 2003-093058/08.

ç Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4257; 763pp; English.

This invention describes a novel composition (a) a first active agent, accomprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and controlly adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenargic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to crackuce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and associated compliance and or surfactant hypoproduction are associated inflammation, allergies and/or bronchoconstriction, vespiratory distance, pulmonary controller pain, cystic fibrosis, allergic rhinitis, pulmonary charactory, pain, cystic fibrosis, allergic rhinitis, pulmonary charactory, emphysema, chronic obstructive pulmorary disease, pulmonary

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transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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i, Shahabuddin S;
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                                                                                                                                                                                                                                                                                                                                                                                                ABD25409 standard; DNA; 20 BP
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                                                                                                                                                                                              Best Local Similarity 100.
Matches 20; Conservative
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Tang L,
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Miller S,
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treating a respiratory, ingo or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lungs or disease or condition such as pulmonary vasoconstriction, inflammation, allergies and/or bronchoconstriction and/or lungs. The confinition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory disease or condition such as pulmonary disease, pulmonary confirmed, pain, cystic fibrosis, allergic rhintis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the chigonucleotides into products that free adenosine into the system content any unwanted effects due to it
     analgesic, hypotensive, immunosuppressive and cytostatic activity, is
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Sequence 20 BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Preα. ... Conservative Similarity 20; Query Match Local Best Loca Matches

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2708 TAAAAAAAAAAAAAAA 2727 요 ઠે

ABD25169 standard; DNA; 20 ABD25169 RESULT

BP. (first entry) 29-JUL-2004 ABD25169; 

AI041482-derived oligonucleotide SEQ ID 4181.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic fullitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, S. Sandrasagra A, ,, Shahabuddin Li Y, Sar Tang L, Nyce JW, Miller S,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4181; 763pp; English

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comprising oligonucleotides, effective for alleviating active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine eansitivity, levels of adenosine (A) or (A) receptors, bronchoconstriction, respiratory tract inflammation, allergies and can ensity adenosine encoding or regulating coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonts. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary viscoconstriction, and or inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary cramplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that the tere edenosine into the sayetem e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to transplantation rejection, election the serves to prevent any unw
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invention describes a novel composition (a) a first active agent,
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100.0%; Pred. No. 6.8
:ive 0; Mismatches
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nes 20; Conservative
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Matches
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Comprising oligonucleotides, effective for alleviating betaform describes a novel composition, alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and control adentiform or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antisathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or radiability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cyllammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung or a disease or condition such as pulmonary visconstriction, and seases or condition such as pulmonary disease, pulmonary cransplantation rejection, pulmonary infections, brain, heart, kidney, etc. tissue environment and thereby, to the oligonucleotides into products that free adenosine to be e.g., lung, brain, heart, kidney, etc. tissue environment and thereby, to the oligonucleotides into product that the encode of the oligonucleotides into the erget encode of the oligonucleotides into the e Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. This invention describes a novel composition (a) a first active agent S Gape oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and Pabalan J, Aguilar D; ö Pharmaceutical composition for treating asthma, has antisense Length 20; 0; Indels Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 0.7%; Score 20; DB 1; Le 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; AI122689-derived oligonucleotide SEQ ID 3807. Katz E, Claim 15; SEQ ID NO 4483; 763pp; English Sandrasagra A, Ka L, Shahabuddin S; 2709 AAAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAAA 20 BP. ABD24795 standard; DNA; 20 (first entry) Best Local Similarity 100. Matches 20; Conservative bronchodilating agent. WPI; 2003-093058/08 29-JUL-2004 Nyce JW, I Miller S, ABD24795; Query Match RESULT 538 ABD24795 ઢ

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This lime and describes a novel composition of a lirst active agent, comprising oligonuclectides, effective for alleviating oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and contending adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction also describes a kit, that composition or cancer and can be anti-sense to the corresponding mana. The invention also describes a kit, that composition of the invention has antiallargic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a nalgesic, hypotensive, immunosuppressive and cytostatic activity, is a creating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung with a disease or condition such as pulmonary viscocrated with a disease or condition such as pulmonary disease, pulmonary contransion, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary diseases, pulmonary contransion, emphysema, chronic obstructions, and emphysema, chronic obstructions, and emphysema, chronic obstructions, and emphysema, chronic obstructions, burnontinis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakpann of the anti-sense oligos corresponding to thymiding present in the target RNA serves to prevent the breakpann of the propriet of thymiding present in the target RNA serves to prevent the breakpann of the propriet of the propriet of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                             Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                             Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 15; SEQ ID NO 3807; 763pp; English.
                                                                                                                                                                                                                                                                                                                          Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                             23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                24-APR-2001; 2001US-0286036P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-093058/08.
                                                     WO200285309-A2.
  Homo sapiens.
                                                                                                          31-OCT-2002
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                              Gaps
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n 0.7%; Score 20; DB 1; Length 20; Similarity 100.0%; Pred. No. 6.8e+02; 20; Conservative 0; Mismatches 0; Indels
     Length 20;
      Local St. 20;
    Query Match
                             Matches
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ABD25110 standard; DNA; 20 BP (first entry) 29-JUL-2004 ABD25110; ABD25110 

RESULT 539

AI125228-derived oligonucleotide SEQ ID 4122.

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatiory; antiathmatic; ansalgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

t

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

Katz E, Nyce JW, Li Y, Sandrasagra A, Ka Miller S, Tang L, Shahabuddin S; (EPIG-) EPIGENESIS PHARM INC.

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Aguilar

Pabalan J,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4122; 763pp; English.

This invention describes a novel composition (a) a lirst active agent, comprising oligonucleotides, effective for alleviating brunchlond describes and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configurated depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallammatory, antiathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a conferency of a great respiratory, lung or malignant disease. The administered corrected treating a respiratory, lung or malignant disease. The administered corrected composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the carget mRNA or to reduce the amount of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The composition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, the disease pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligon corresponding to the rearget RNA serves to prevent the breakdown of the uniquence and thereby, to thymidines present in the target RNA serves to prevent the breakdown of edge. The reduced adenosine content of the anti-sense of the prevent of the serves to prevent the prevent of the p This invention describes a novel composition (a) a first active agent e.g., lung, brain, heart, kidney, etc, prevent any unwanted effects due to it

Sequence 20 BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;

Gaps . 0 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels 20; Conservative Query Match Best Local Similarity Matches

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating cy required to a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a curating a respiratory, lung or mailgnant disease. The administered correcting a respiratory, lung or mailgnant disease. The administered correction comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and or canner. The rangel canner, and the effect of pain, cystic fibrosis, allergic rhinitis, pulmonary rangel efforms in pain, respiration, respiratory emphysema, chronic obstructive pulmonary disease, pulmonary rangelentsion, emphysema, chronic obstructive pulmonary disease, pulmonary rangelentsion, emphysema
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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Tang L, Shahabuddin S;
2708 TAAAAAAAAAAAAAAA 2727
                                             ABD25934 standard; DNA; 20 BP
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transplantation rejection, pulmonary infections, bronchitis or cancer.

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antifinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                        Gaps
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100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indela
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                                                                                                                         Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA505075-derived oligonucleotide SEQ ID 4947.
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Tang L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                            ABD25935 standard; DNA; 20
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                                                                                                                                                                                                          20; Conservative
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                                                                                                                                                                     Query Match
Best Local Similarity
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treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary hypertension, emphysema, chronic obstructions, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of energy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              prevent any unwanted effects due to it
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels 100.0%; Prec. ... 0.78; 20; Conservative Local Similarity Query Match Matches ò

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Gaps

2709 AAAAAAAAAAAAAAA 2728 1 AAAAAAAAAAAAAAAAA 20 g

RESULT 542

ABD25936 standard; DNA; 20 BP ABD25936

ABD25936;

(first entry) 29-JUL-2004

AA505075-derived oligonucleotide SEQ ID 4948.

Human; antieense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthastic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar Katz E, Tang L, Shahabuddin S; Sandrasagra A, Li Y, Nyce JW, L Miller S, 

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4948; 763pp; English.

This invention describes a novel composition (a) a first active agent,

Computation displantation, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant deplation or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a container agonetic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or tracer polypeptide present in the lungs. The cumposition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung or inflammation, allergies and/or surfactant hypoproduction are associated with a disease or conditions and as pulmonary vasoconstriction, compliation such as pulmonary disease, pulmonary contains and present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system content any unwanted effects due to it comprising oligonucleotides, effective for alleviating

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Preα. ... tive 0; Mismatches Ouery Match
Best Local Similarity 100...
Best Local 20; Conservative

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ò 셤 RESULT 543 ABD21541/c ABD21541;

ABD21541 standard; DNA; 20 BP

(first entry) 29-JUL-2004

S100 calcium binding protein A2-derived oligo SEQ ID 553.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disterses syndrome; allergic fuinitis; pulmonary vasoconstriction; respiratory distratory distratory distratory distratory distratory distratory distratory distratory pulmonary dispasse; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. 

Homo sapiens.

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WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Katz E, Pabalan J, Nyce JW, Li Y, Sandrasagra A,

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comprising oligonucleotides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cuffactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a corresponding a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to craduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung contains and or bronchic present in the larges or condition such as pulmonary vasoconstriction, and or bronch produce the anti-sense or condition such as pulmonary vasoconstriction, conficients or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the inflammation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine the breakdown of the oligonucleotides into products that free edenosine the products that free edenosine the products due to it the envisonment and thereby, to the oligonucleotides into products that free edenosine to it
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                                                                                                                                                                                                                                                                                   This invention describes a novel composition (a) a first active agent,
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                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 553; 763pp; English.
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Tang L, Shahabuddin S;
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Best Local Similarity 100...
Conservative
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                                                                                                                                                                                     bronchodilating agent
                                                  WPI; 2003-093058/08
     Miller S,
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for illammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating cyfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) analgesic, hypotensive, immunosuppressive and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antianflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition, and/or bronchoconstriction and/or lung or reduce the amount of target polypeptide present in the lungs. The cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, contained the intension, content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the saster the oligos corresponding to the oligonucleotides into products that free adenosine into the saster the suggest of the oligonucleotides into products that free adenosine and therefore the submonst.
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                                                                                                                                                                                                                                                                                                                                              oligonucleotide containing less percentage of adenosine, targeted toncleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                        Pabalan J, Aguilar D;
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                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
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                                                                                                                                                                                                                           Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 15; SEQ ID NO 4683; 763pp; English
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Tang L, Shahabuddin S;
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                                                                                                                                          24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                 (EPIG-) EPIGENESIS PHARM INC
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Les 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                  bronchodilating agent
                                                                                                                                                                                                                                                                                       WPI; 2003-093058/08.
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                 WO200285309-A2
                                                          31-OCT-2002
                                                                                                                                                                                                                                              Miller S,
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Ното варіеля

AA278764-derived oligonucleotide SEQ ID 5892.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory distantery distress syndrome; allergic rhinitis; pulmonary vasoconstriction; respiratory distructive pulmonary disease; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002,

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

ä Aguilar Pabalan J, Katz E, Tang L, Shahabuddin S; Sandrasagra A, Li Y, Miller S, Nyce JW,

WPI; 2003-093058/08.

S oligonuclectide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and Pharmaceutical composition for treating asthma, has antisense bronchodilating agent.

Claim 15; SEQ ID NO 5892; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating control of the composition, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and ceducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating cypression of a target polypeptide associated with lung airway or lung dysfunction also describes a kit, that comprises: (a) a delivery of the invention also describes a kit, that comprises: (a) a delivery of the invention as carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition or availability, or to increase the administered to reduce the production or availability, or to increase the administered to reduce the production or availability, or to increase the administered to reduce the production or availability, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and or cancer. The inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary the radvord adanced to the radvord distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary the radvord adanced to the radvord distress syndrome, pain, cystic fibrosis, allergic reduced the adaptory of the transplantation, emphysema, chronic observations, productions, produced reduced the advording or condition or availed to the condition and the The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of invention describes a novel composition (a) a first active agent,

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it

Gaps .; 0 Length 20; Indels 6.8e+02; DB 1; 0.7%; Score 20; DB 100.0%; Pred. No. 6.8 :ive 0; Mismatches 0./\* Dest Local Similarity 100.0 Matches 20, Conservative

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1 AAAAAAAAAAAAAAA 20 

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RESULT 546

ВÞ ABD24850 standard; DNA; 20

ABD24850;

(first entry) 29-JUL-2004

AI092623-derived oligonucleotide SEQ ID 3862.

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diserses syndrome; allergic rhintis; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

(EPIG-) EPIGENESIS PHARM INC.

24-APR-2001; 2001US-0286036P.

Aguilar D; Katz E, Pabalan J, Nyce JW, Li Y, Sandrasagra A, Ka Miller S, Tang L, Shahabuddin S; 

WPI; 2003-093058/08.

S Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

Claim 15; SEQ ID NO 3862; 763pp; English.

The invention also describes that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a nalgesic, hypotensive, immunosuppressive and cytostatic activity, is a cheat addrenesgic agonist. The composition is useful for preventing or treating a respiratory, lung or mailgnant disease. The administered composition comprises oligo and is administered to reduce the production or treating a respiratory, lung or mailgnant disease. The administered corposition of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. This invention describes a novel composition (a) a first active agent,

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition is useful for preventing or beta-adrenergic agonist. The composition is useful for preventing or
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thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, it prevent any unwanted effects due to it
                                                                                                                                                                      Gaps
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                                                                                                                                 DB 1; Length 20; 6.8e+02;
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                                                                                           BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI125651-derived oligonucleotide SEQ ID 4544.
                                                                                                                               0.7%; Score 20; DB
100.0%; Pred. No. 6.8
tive 0; Mismatches
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S;
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,, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                              ABD25532 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                    Best Local Similarity 100.
Matches 20; Conservative
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Tang L,
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                                                                                             Sequence 20
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treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung to inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction.

Cinflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it Human, antisense, bronchoconstriction, allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; crostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating Gaps oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and ä ö Aguilar Pharmaceutical composition for treating asthma, has antisense Score 20; DB 1; Length 20; Pred. No. 6.8e+02; Indels Pabalan J, Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; AI128305-derived oligonucleotide SEQ ID 4058. 100.0%; Prea. ... Katz E, Claim 15; SEQ ID NO 4058; 763pp; English. ŝ 2709 AAAAAAAAAAAAAAAA 2728 20 Sandrasagra A, L, Shahabuddin 1 AAAAAAAAAAAAAAAAAA BP. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC ABD25046 standard; DNA; 20 (first entry) 20; Conservative bronchodilating agent WPI; 2003-093058/08. Li Y, Sar Tang L, Local Similarity WO200285309-A2 29-JUL-2004 Homo sapiens. 31-OCT-2002. Miller S, ABD25046; Nyce JW, Query Match Matches ABD2504 ID ABI XX 硆 ò

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bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung definition or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialflammatory, antiaethmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The production and/or surface and in administrian and in administrial and inflammation and an analyses of the amount of target polypeptide present in the lungs. The administrial and an analyse of the amount of target polypeptide present in the lungs. The analyse and an administrial and an administ
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Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

. 0 0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Pred. No. 6.8 ive 0; Mismatches 2709 AAAAAAAAAAAAAAA 2728 Query Match
Best Local Similarity 100."
Matches 20, Conservative à

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Gaps

1 AAAAAAAAAAAAAAA 20

ABD25044 standard; DNA; 20 BP 29-JUL-2004 (first entry) ABD25044; RESULT 549 ABD25044 

AI128305-derived oligonucleotide SEQ ID 4056.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4056; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating by the comprising oligonucleotides, effective for alleviating constriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the variet papers of the administered containers are an experienced or composition or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of tearget polypeptide present in the lungs. The cultinammation, allergies and/or surfactant hypoproduction are associated confinammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction, respiratory confisteres syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary confistered adenosine content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos corresponding to the holison relection, pulmonary infections, planers into the anti-sense oligos corresponding to the holison of the partice and planers into the parget RNA serves to prevent the breaders. kidney, etc, tissue environment and thereby, to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, prevent any unwanted effects due to it 

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; 0; Indels 100.0%; Preα. ... tive 0; Mismatches Local Similarity 100. 1es 20; Conservative Query Match Best Loca Matches

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ABD25111

ABD25111 standard; DNA; 20 BP ABD25111; 

All25228-derived oligonucleotide SEQ ID 4123.

(first entry)

29-JUL-2004

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fihitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant deplation or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating creating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a container, immunosuppressive and cytostatic activity, is a container, immunosuppressive and cytostatic activity, is a container, and in a useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligons corresponding to the villagence or condition usuch as pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligonscleeding into products that the eastern in the target RNA serves to prevent the breakdown of the oligonucleotid
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                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                        Pabalan J, Aguilar D;
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100.0%; Pred. No. 6.8e+02;
vative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                        Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 4123; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g., lung, brain, heart, kidney, etc, prevent any unwanted effects due to it
                                                                                                                                                                                                                      Nyce JW, Li Y, Sandrasagra A, Ka
Miller S, Tang L, Shahabuddin S;
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ADZ20571 standard; DNA; 20 BP.
                                                                                        23-APR-2002; 2002WO-US013143.
                                                                                                                                  24-APR-2001; 2001US-0286036P
                                                                                                                                                                            (EPIG-) EPIGENESIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                       bronchodilating agent
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WO200285309-A2
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Local Sim-
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Gene expression detection related oligo, SEQ ID 3.

(first entry)

16-JUN-2005

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The invention relates to a novel method for amplifying a nucleic acid.

The method involves: synthesizing a DNA fragment having a first consensus sequence, which is not contained in the 5' terminal of the nucleic acid; adding a homopolymer to the 3' and using a terminal deoxynucleotide transferase; synthesizing a DNA fragment having a second consensus cransferase; synthesizing a DNA fragment having a second consensus containing PCR using the first and second primers to the homopolymer; and performing PCR using the first and second primers. The invention further comprises: a method for preparing a labeled nucleic acid, which involves the steps of the novel method above in which the PCR is performed using a label or primers; a nucleic acid detection system containing labeled nucleic acid, nucleic acid probe and immobilized solid-phase support body is a gene-expression monitoring system, comprising a solid-phase support or a nucleic acid probe and labeled nucleic acid; a kit for amplifying comprising the minal deoxynucleotide; DNA polymerase, first and second consensus consensus sequences; and a kit for labeling conclosing the components of the kit and labeled conclosing the components of the kit and labeled conclosing nucleic acids, comprising the components of the kit and labeled concloses. This polynucleotide sequence represents an oligonucleotide conclosion in the method for detecting nucleic acids expressed in a sample of
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                DNA amplification; DNA detection; gene expression;
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                                                                                                                                                                                                                                                           (TOYM ) TOYOBO KK.
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                                                          Synthetic.
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Detecting nucleic acid used for e.g. diagnosis of diseases, forensics and DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
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                                                                                                                                                                                                                                                                         The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                        Storhoff JJ, Elghanian R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
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nanoparticle-oligonucleotide conjugate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                            Example 18; SEQ ID NO 55; 130pp; English
                                                        Mucic RC,
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97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0200161P.
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                                                        Letsinger RL,
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                   (NANO-) NANOSPHERE INC.
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les 20; Conserv
                                                                                                                                                                                                         oligonucleotides.
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26-APR-2000;
26-JUN-2000;
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                                                        Mirkin CA,
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                        Elghanian R;
                                                                                                                                                                                                                                                                                                            Detecting nucleic acid used for e.g. diagnosis of diseases, for DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
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nanoparticle-oligonucleotide conjugate.
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                                                29-JUL-1996; 96US-0031809P.
21-JUL-1997; 97WG-US012783.
29-JUN-1999; 99US-00344667.
25-JUN-1999; 99US-00344667.
26-JUN-2000; 2000US-0200161P.
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               10-OCT-2001; 2001US-00973638
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Matches 20; Conservative
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26-APR-2000;
26-JUN-2000;
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21-JUL-1997;
29-JAN-1999;
                                              29-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                           The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking original oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid amplification; RNA transcription; RNA polymerase; ss; T7.
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DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide sequence of a dA20 oligonucleotide.
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                                                                             Example 18; SEQ ID NO 55; 130pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI34492 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-035466/03.
                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
nes 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kamme FC, Zhu JY;
                                        oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003102243-A1.
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Best Local 5
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                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis of sample molecules such as DNA fragment, by using micro channel to form laminar flow of specimen molecule-containing solution and complex forming molecule containing solution.
                                                                                                                                                                                                                                                                                                                                                                                                                            laminar flow; micro channel; complex; selectively promoted; fluorescence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                  Gaps
template derived production of RNA in the transcription reaction. The present sequence represents an oligonucleotide used to exemplify RNA transcription in the presence of single- and double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
:ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                               Molecule analysing microchannel method related probe #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligo used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shimizu H, Miyazaki M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
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                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2002; 2002JP-00211462.
                                                                                                                                                                                                                                                                                                    ADI47212 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                    20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-180318/17.
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Best Local Similarity
                                                                                                                       Query Match
Best Local Similarity
                                                    oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yamashita K,
Yamaguchi Y;
                                                                                                                                                                                                                                                                                                                                                                   22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe;
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Matches
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Indels

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20; Conservative

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a peptide nucleic acid analogue. OC is useful as a prodrug, useful in diagnostics, therapeutics and as research reagents and kits. OC is useful for preventing or delaying infection, inflammation or tumour formation in organisms. The present sequence represents an oligonucleoride used in the
                                                                                                                                                                                                                                                                                                                                /note= "Optionally conjugated with spermine, polyethylenimine (PEI) 600 or PEI 1200, tetraethylenpentamine. Also optionally 5'-protected with DMT."
                                                                                                                                                                         88; Antimicrobial; Antiinflammatory; Cytostatic; prodrug; infection; inflammation; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polyalkyleneamine-containing oligomeric compound useful for preventing or delaying infection, inflammation or tumor formation in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                         Polyalkyleneamine-conjugated oligonucleotide #1.
                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maier MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 22; 37pp; English
                                                                                                                                                                                                                                                                                                                          base= OTHER
                              ADJ51142 standard; DNA; 20 BP.
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                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                           *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MANO/) MANOHARAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GUZA/) GUZAEV A P. (MAIE/) MAIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                     JS2004019000-A1
                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Manoharan M,
                                                                                                 06-MAY-2004
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                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organisms.
                                                              ADJ51142;
           ADJ51142/
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0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;

Query Match Best Local Similarity

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having at least two portions comprising contacting the nucleic acid with at least two types of nanoparticles, such as gold, having attached oligonuclectides and observing a detectable change brought about by hybridisation of the oligonuclectides on the nanoparticles with the nucleic acid. The method of the invention may be useful for detecting a nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene associated with a disease, a fungal DNA, synthetic DNA or RNA, structurally modified natural or synthetic DNA or RNA or a product of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of nucleic acid, e.g. viral RNA or DNA, comprises contacting nucleic acid with different types of nanoparticles having attached oligonuclectides and observing detectable change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for detecting a nucleic acid
                                                                                                                                                                                                                                     nanoparticle, gold, disease, forensic, paternity testing, cell line authentication, gene therapy, ss, gold colloid conjugate.
                                                                                                                                                                                                      Oligo related to thiol oligo-gold colloid conjugate probe SEQ 70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 24; SEQ ID NO 70; 206pp; English.
2709 AAAAAAAAAAAAAAAA 2728
                                AAAAAAAAAAAAAAAA
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                                                                                                          ADI32920 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                               97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0176409P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Taton TA, Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                08-OCT-2002; 2002US-00266983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-DEC-2001; 2001US-00008978
                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                  US2003207296-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PARK/) PARK S. (TATO/) TATON T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUN-1999;
13-JAN-2000;
28-MAR-2000;
26-APR-2000;
                                                                                                                                                                      06-MAY-2004
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                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                              RESULT 558
                                                                                             ADI32920
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polymerase chain reaction amplification. The detected nucleic acid may be utilised for diagnosis of disease, sequencing of nucleic acids, forensics, paternity testing, cell line authentication and monitoring gene therapy. The method for detecting the nucleic acids is based on observing a colour change with the naked eye and is cheap, fast, simple, and robust, requiring no specialised or expensive equipment. The current sequence is that of the oligonucleotide which is related to a thiolmodified oligonucleotide-gold conjugate probe of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detection of nucleic acid, e.g. viral RNA or DNA, comprises contacting nucleic acid with different types of nanoparticles having attached
                                                                                                                                                                                                                                                                                                                                                                      - SEQ 55
                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                               nanoparticle, gold, disease, forensic, paternity testing, cell line authentication, gene therapy, ss; gold colloid conjugate;
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0
                                                                                                                                                                                                                                                                                                                                                                      Synthetic thiol-modified oligo-gold colloid conjugate probe
                                                                                                                                            0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                     Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                   2000US-0213906P.
2000US-022461P.
2000US-025418P.
2000US-0255238P.
2000US-0255238P.
2001US-0076050.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0176409P.
2000US-0192699P.
2000US-0200161P.
2000US-00603830.
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99US-00240755.
99US-00344667.
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2001US-0282640P.
2001US-00927777.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-OCT-2002; 2002US-00266983
                                                                                                                                                                                                                                                                                           ADI32905 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                          Conservative
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                                                                                                                                                           Local Similarity
ses 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taton TA,
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11-AUG-2000; 2
08-DEC-2000; 2
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11-DEC-2000;
11-DEC-2000;
12-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                           probe.
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Matches
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ID ADI3
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The invention relates to a novel method for detecting a nucleic acid having at least two portions comprising contacting the nucleic acid with at least two portions comprising contacting the nucleic acid with a least two types of nanoparticles, such as gold, having attached oligonuclectides and observing a detectable change brought about by hyridisation of the oligonuclectides on the nanoparticles with the nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene associated with a disease, a fungal DNA, synthetic DNA or RNA, cuturally modified natural or synthetic DNA or RNA or a product of a polymerase chain reaction amplification. The detected nucleic acid may be utilised for disgnosis of disease, sequencing of nucleic acids, forenaics, paternity testing, cell line authentication and monitoring gene therapy. The method for detecting the nucleic acids is based on observing a colour change with the naked eye and is cheap, fast, simple, and robust, requiring no specialised or expensive equipment. The current cand robust, requiring no specialised or expensive equipment. The current cand colour change with the invention which is linked via a thiol
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gape
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oligonucleotides and observing detectable change brought about by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Phosphorothioate backbone; sulphurised oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                       Example 18; SEQ ID NO 55; 206pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sulphurised oligonucleotide #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-JAN-2000; 2000US-00481486.
10-JAN-2001; 2001WO-US000715.
                                                                                                                                                                                                                                                                                                                                                                                         group to a gold nanoparticle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-DEC-2002; 2002US-00181200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADK69880 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004 (first entry)
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CHERUVALLATH Z S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.7
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cole DL, Ravikumar VT,
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003212267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
                         hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified.
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(CHER/)
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Matches
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                                                                                                                                                                                                                                                                              phosphite intermediate, and oxidising the phosphite intermediate with an acetyl disulfide in an acetonitrile for a time to effect conversion of the phosphite intermediate to phosphorchicate. The invented method achieves high yields and greater efficiency. The present sequence is sulphurised oligonucleotide used in the exemplification of the invention.
                                      to
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Preparation of phosphorothioate oligonucleotides involves oxidizing phosphite intermediate with acetyl disulfide in acetonitrile for time effect conversion of phosphite intermediate to phosphorothioate.
                                                                                                                                                                 The invention relates to phosphorothioate oligomucleotides having nucleoside with 240 modification are prepared by phosphitylating 5'-hydroxyl of a nucleic acid moiety having a nucleoside with 2' modification in an acetonitrile containing solvent mixture to form a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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/note= "Phosphorothioate backbone; 2'-O-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to phosphorothioate oligonucleotides having nucleoside with 240 modification are prepared by phosphitylating 5'-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preparation of phosphorothioate oligonucleotides involves oxidizing phosphite intermediate with acetyl disulfide in acetonitrile for tin effect conversion of phosphite intermediate to phosphorothioate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 20; DB 1; Le
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. No.
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                                                                                                               Example 12; SEQ ID NO 10; 8pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JAN-2000; 2000US-00481486.
10-JAN-2001; 2001WO-US000715.
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/*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COLE D L.
RAVIKUMAR V T.
CHERUVALLATH Z S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20; Conservative
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound pecifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, classific meuropathy, trigeminal neuropathy, arthritic pain, acute pain, cheadache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with chuman Nav1.3 expression, the oligonucleotides are designed to target clifferent regions of the human Nav1.3 RNA.
hydroxyl of a nucleic acid moiety having a nucleoside with 2' modification in an acetonitrile containing solvent mixture to form a phosphite intermediate; and oxidising the phosphite intermediate with an acetyl disulfide in an acetonitrile for a time to effect conversion of the phosphite intermediate to phosphorchhioate. The invented method achieves high yields and greater efficiency. The present sequence is sulphurised oligonucleotide used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound targeted to a nucleic acid molecule encoding abar1.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chimeric phosphorothioate oligonucleotide to target Nav1.3 #2303.
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                                                                                                                                                                                                       0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                             ВР
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                                                                                                                                                                                                                                               20; Conservative
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                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound concerned and composition are useful for treating a disease or condition associated with Navl.3. e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, caspeciated with Navl.3. e.g. pain including but not limited to associated with Navl.3 e.g. pain including but not limited to pain, compound triggation headache, cluster headache, mild-to-moderate cheadache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorchioate oligonucleotide with 2.MoE wings and a deoxy gap. Used during the antisense inhibition of thuman Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothicate oligonucleotide to target Nav1.3 #2223.
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Pred. No. 6.8e+02;
                   Length 20;
                                                                        Indels
DB 1; Leads 1. 6.8e+02;
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                   0.7%; Score 20; DB
100.0%; Pred. No. 6.8
ive 0; Mismatches
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                                                                                                                                                               ADK74889 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                           Conservative
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Best Local Similarity
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           Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W02004016754-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                              ADK74889;
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                                                                                                                                                                                                                                                                    RESULT
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The present invention relates to a method (M1) for detecting and/or isolating a nucleic acid having a homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) comprises treating a sample containing nucleic acid compounds with an locked nucleic acid (LNA) oligonucleotide (LO) to thereby detect and/or isolate a nucleic acid having the homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) is useful for detecting and isolating nucleic acids released from a lysed complex biological mixture comprising nucleic acids. The present sequence is a LNA oligomer, used to illustrate the
                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "Optionally biotinylated or 5' AQ2-HEG3, where AQ
is anthraquinone and HEG is hexa-ethylene glycol"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting and/or isolating nucleic acid molecule having homopolymeric sequence or repetitive element or conserved nucleotide sequence involves treating sample containing nucleic acid compounds with locked nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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100.0%; Pred. No. 6.8e+02;
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                                                                                                                         Detection, isolation; locked nucleic acid; LNA; ss.
                                                                                                                                                                                                                                    /note= "Optionally LNA nucleotides"
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                                                                                                                                                                                 location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 22; Page 51; 104pp; English.
                                                                                                                                                                                                                         /mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
20-JUN-2003; 2003WO-IB006354
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ADM13992/
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Conservative

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RESULT 564

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytosetatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antidifiammatory; neuroprotective; nautionifiammatory; neuroprotective; vasotropic; ophthalmological; neuroprotective; adisease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                         Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:179.
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/note= "2'-0-methocyethyls"
16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mod_base= OTHER
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(first entry)
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                         sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human meGES-1 gene is located on chromosome 9, more specifically to days. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, in cells or tissues; and (3) a method of inhibiting the expression of mPGES-1, in cells or tissues; and (3) a method of treating an animal natisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 er, inflammation, Alzheimer's condition associated with mPGES-1 er, inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.

Claim 4; SEQ ID NO 179; 132pp; English.

ischemia

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; implibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; immunomodulatory; cardiant; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds,
                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
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                                                                                                                                                                                                                                            Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:181.
           Length 20;
                                   0; Indels
       DB 1; Ler
. 6.8e+02;
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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16. .20
                                   Mismatches
          Score 20;
Pred. No.
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                                                               2728
0.7%; Sc.
100.0%; Pred
0; M
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/mod_base= OTHER
                                                                                        20 AAAAAAAAAAAAAAA 1
                                                             2709 AAAAAAAAAAAAAAAAA
                                                                                                                                                             BP
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                                                                                                                                                .994/c
ADM13994 standard; DNA; 20
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                                     Conservative
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*tag=
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                     Similarity
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        Query Match
Best Local Simi:
Matches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                 RESULT 566
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having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MPGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; pPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nationiflammatory; neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                          chritis, diabetes, cancer, ischaemia or reperfusion injury, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:186.
                                                                                                                                                                                                                                                                                                                  0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         residues are 5-methylcytidines
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/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                            arthritis,
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modified_base
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-2004
                                                                                                                                                                                                                                               ophthalmic,
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ADM13999/c
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to ogga4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition aspociated with mPGES-1 e.g., inflammation, Altheimer's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                Claim 4; SEQ ID NO 186; 132pp; English.
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                                                            WPI; 2004-305094/28.
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ADM14008/c
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                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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100.0%; Pred. No. 6.8e+02;
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                           /mod_base= OTHER
/note= "2'-O-methoxyethyls"
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tive 0; Mismatches
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                                                             *tag=
                                                                                                                                                                                                                                                                                     (PHAA ) PHARMACIA CORP
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Matches 20; Conserv
                                                                                                                                    WO2004028458-A2
                                       modified base
                                                                                                                                                                        08-APR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ischemia.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically too 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically thybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal artisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamacroy, neuroprotective, nootropic, antidiamacroy, neuroprotective, nootropic, antidiamacroy, neuroprotective, nootropic, antidiamacroy, neuroprotective, and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound con all be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9d34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidlabetic, immunomodulator, cardiant, neuroprotective, antidlabetic, jamunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and cardiavascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or disease o
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/note= "phosphorothioate linkages and all cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:184.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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/note= "2'-0-methocyethyls"
16. .20
                          Claim 4; SEQ ID NO 338; 132pp; English
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Best Local Similarity 100.0
Matches 20; Conservative
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Synthetic.
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The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mBCBS-1). The human mPGBS-1 gene is located on chromosome 9, more specifically to of 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGBS-1, which specifically whybridise with the nucleic acid encoding mPGES-1 in cells or tissues; and (3) a method of treating an animal charing a disease or condition associated with mPGBS-1. MPGBS-1 and antidiabetic, immunomodulator, cardiant, neuroprotective, cardiant, neuroprotective, antidiabetic, vasotropic, antidiabetic, immunomodulator, and cardiovascular activities, and can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound to phthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGBS-1 inhibitors and in gene therapy. The antisense compound condition associated with mPGBS-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or condition associated with mPGBS-1 e.g., inflammation, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory, neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; repertusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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Pred. No. 6.8e+02;
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ive 0; Mismatches
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                                        25-SEP-2003; 2003WO-US030374.
                                                                              25-SEP-2002; 2002US-0413549P.
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                                                                                                                     (PHAA ) PHARMACIA CORP
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Synthetic.
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  08-APR-2004.
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                                                                                                                                                                                                                                                                                                            ischemia.
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Location/Qualifiers

Key

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mFGES-1; mFGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:205.

/mod\_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

/mod\_base= OTHER

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/\*tag=

modified\_base

Location/Qualifiers

Key modified\_base

Homo sapiens

Synthetic.

\*tag=

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mEGES-1). The human meGES-1 gene is located on chromosome 9, more specifically to oliginal properties and meGES-1, which specifically hybridise with the nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding in thibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisance oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1. Grimmation, Althebiaer's condition associated with mPGES-1 e.g., inflammation, Althebiaer's condition associated with mPGES-1 e.g., inflammating a disease or condition associated with mPGES-1 e.g., inflammation, Althebiaer's condition associated with mPGES-1 e.g., inflammation, Althebiaer, e.g., e.g.
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                                        /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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16. .20
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Best Local Similarity
Matches 20; Conserv
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modified_base
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to human mpGES-1 gene is located on chromosome 9, more specifically to having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1 which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal comparing a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, nootropic, antistriitie, vasotropic, antidiamaticit, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound to the used as mPGES-1 inhibitors and in gene therapy. The antisense compound con the used as mPGES-1 inhibitors and in gene therapy. The antisense compound to condition associated with mPGES-1 e.g., inflammation, Altheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or condition associated with mPGES-1 e.g., inflammation, injury, or condition associated with activities condition and conditions and condition and conditions and conditins and conditions and conditions and conditions and conditions an
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'note= "2'-0-methocyethyls"
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AAAAAAAAAAAAAAAAAAA

20

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ADM14018 standard; DNA; 20 BP

RESULT 574 ADM14018, (first entry)

01-JUL-2004

ADM14018;

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08-APR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gierse JK;
                                                                                                                                                                                                                                                                                                     ADM14257:
                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                           RESULT 576
                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                       ADM14257,
         임
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                                                                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nationflammatory; neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                    Gaps
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                                                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:275.
                   Indels
       Pred. No. 6.8e+02;
Mismatches 0;
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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                                     2709 AAAAAAAAAAAAAAAA 2728
100.0%; Prr
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                                                        20 AAAAAAAAAAAAAAAAAA
                                                                                                           ADM14088 standard; DNA; 20 BP
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                                                                                                                                                    (first entry)
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                  Conservative
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       Best Local Similarity
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modified_base
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                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                               ADM14088;
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                 Matches
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                   mpGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1. Chimeric antisense oligonucleotides and antisense compounds have cytostatic, antistabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1. e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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inhibits its expression; (2) a method of inhibiting the expression of
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                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20; DB 1; Lot
Pred. No. 6.8e+02;
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/note= "2'-0-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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ses 20; Conserva
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Seguence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "2'-O-methocyethyls"
                                                                                                                                                              100.0%; Pred. ...
                                       Claim 4; SEQ ID NO 444; 132pp; English.
                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                   ADM14000 standard; DNA; 20 BP
                                                                                                                                                            0.7%;
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                                                                                                                                                        Query Match
Best Local Similarity 100.0
....hes 20; Conservative
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modified_base
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Synthetic.
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                              ischemia.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The human meges-1 gene is located on chromosome 9, more specifically to open is located on chromosome 9, more specifically to open is located on chromosome 9, more specifically to open is sequence comprising 8-30 bp targeted to a nucleic acid encoding mpCES-1, which specifically Mybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPCES-1 in cells or tissues; and (3) a method of treating an animal converse or condition associated with mPGES-1. MPCES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, cantidiabetic, immunomodulator, cardiant, neuroprotective, neotropic, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPCES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPCES-1 e.g., inflammation, Alzheimer's condition associated with mPCES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                /note= "2'-0-methoxyethyls"
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                                                 /mod_base= OTHER
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                              *tag=
                                                                                                                                                                                                                                                                                                                                                                                 (PHAA ) PHARMACIA CORP
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modified base
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                                                                                                                                                                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to operate the present invention also describes: (1) antisense compounds, earlies a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or disparance and with mPGES-1 e.g. inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                 encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                            antisense compound, having a sequence targeted to a nucleic acid
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
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                              WPI; 2004-305094/28
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Gaps ö

7 10:41:34 2006

Tue Nov

ADM14014/c

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to oliginally to adving a sequence comprising 8-30 bp targeted to a nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1, which specifically hybridise with mPGES-1. MPGES-1 and inhibits of tissues, and oliginally an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and aradiant, neuroprotective, antinflammatory, neuroprotective, antinflammatory, neuroprotective, antisense compound second as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or disease or disease or condition associated with mPGES-1 e-g., inflammation, Alzheimer's disease or dise
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                 /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "2'-O-methocyethyls"
[6. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "2'-0-methoxyethyls"
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                                                                                                                                                            Location/Qualifiers
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                                                                               Homo sapiens.
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Query Match 0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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2709 AAAAAAAAAAAAAAAA 2728
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20 AAAAAAAAAAAAAAA 1
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승 원

The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to ogg4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antistabletic, immunomodulator, cardiant, neuroprotective, antistiamatory, neuroprotective, nootropic, antistiamatory, neuroprotective, and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase; imPGBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytoatatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antidifammatory; neuroprotective; nationifammatory; immunomodulatory; cardiant; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines" Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:201. /\*tag= a /mod\_base= OTHER /note= "2'-O-methocyethyls" 16. .20 /note= "2'-0-methoxyethyls" Claim 4; SEQ ID NO 201; 132pp; English. Location/Qualifiers /mod base= OTHER OTHER ADM14014 standard; DNA; 20 BP. 25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P. /\*tag= c /mod\_base= (first entry) Д ...20 \*tag= (PHAA ) PHARMACIA CORP. WPI; 2004-305094/28. WO2004028458-A2 modified\_base modified base modified base sapiens 01-JUL-2004 08-APR-2004. Homo sapie Synthetic. Gierse JK; ADM14014; ischemia. Key 

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condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, ophthalmic, immunological, cardiovascular or neurological disorder.
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                       . Match 0.7%; Score 20; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 6.8e+02; les 20; Conservative 0; Mismatches 0; Indels
                                                             Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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/note= "2'-0-methocyethyls"
16. .20
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Matches
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The thuman mPGES-1 gene is located on chromosome 9, more specifically to having a sequence comprising 8-30 bp targeted to antisense compounds, chaving a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding compounds its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal compounds a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, cophthalmological, immunomodulatory and cardiovascular activities, and can ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPGES-1, inflammation, alsease or condition associated with mPGES-1, inflammation, alsease or condition associated with mPGES-1 and can composition for treating a disease or condition associated with mPGES-1 by inflammation, alsease or condition associated with mPGES-1 e.m. included the antisense compound
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                                                                                                                                                                                                                                                                                                                                    disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indels
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16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; neuroprotective; adisease; arthritis; diabetes; cancer; ischaemis, Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0;
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25-SEP-2003; 2003WO-US030374.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 0343.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and inhibiting an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiavascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or discontine and inflammation, Alzheimer's
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/note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
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Pred. No. 6.8e+02;
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16. .20
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antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alfanimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se. The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGBS-1). The human mPGBS-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGBS-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGBS-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGBS-1. MPGES-1 chimeric New antieense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or /mod\_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines" Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; /note= "2'-0-methoxyethyls" note= "2'-0-methocyethyls" Claim 4; SEQ ID NO 192; 132pp; English Location/Qualifiers l. .5 /\*tag= a /mod\_base= OTHER /mod base= OTHER 25-SEP-2003; 2003WO-US030374 25-SEP-2002; 2002US-0413549P. Ω 16. .20 /\*tag= (PHAA ) PHARMACIA CORP WPI; 2004-305094/28. WO2004028458-A2 Key modified\_base modified base modified base Homo sapiens. Synthetic. 08-APR-2004

ö Query Match 0.7%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0; Indels

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Gaps

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mCeBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase; imCeBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antidinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                            Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:182
2709 AAAAAAAAAAAAAAAA 2728
                   20 AAAAAAAAAAAAAAAAAA 1
                                                                     ADM13995 standard; DNA; 20
                                                                                                           01-JUL-2004 (first entry)
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/note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"  $% \left\{ \frac{1}{2}\right\} =\frac{1}{2}$ /mod\_base= OTHER /note= "2'-0-methocyethyls" 16. .20  $not\overline{e} = "2' - 0 - methoxyethyls"$ Location/Qualifiers /mod\_base= OTHER base= OTHER Д ø υ /\*tag= Key modified base modified base modified\_base

sapiens

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25-SEP-2003; 2003WO-US030374.

25-SEP-2002; 2002US-0413549P

(PHAA ) PHARMACIA CORP

Gierse JK;

WPI; 2004-305094/28

antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or encoding mPGESischemia.

Claim 4; SEQ ID NO 182; 132pp; English.

The present sequence represents a chimeric antisense oligonucleotide targeted to human medics.

1. The human medics. Jean is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding medges. I, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 and method of inhibiting an animal

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having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonuclectides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, optication, antiarthritic, vasotropic, antiarthritic, ussotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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00.0%; Pred. No. 6.8e+02;
ve 0; Mismatches 0; Indels
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to gene is located on chromosome 9, more specifically to gay4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal mpGES-1 in cells or tissues; and (3) a method of treating an animal cartisense oligonucleotides and antisense compounds have cytostatic, antidabetic, immunomodulator, cardiant, neuroprotective, cardiant, neuroprotective, neotropic, antistense oligonucleotides and antisense compounds completed as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used as mpGES-1 inhibitors and in gene therapy. Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's diabetes, cancer, isohemana or repertusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. ô chimeric; antisense oligonucleotide; phosphorothioate; human; microscomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microscomal prostaglandin E2 synthase; implibitor; cytostatic; antidiabetic; immunoomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunoomodulatory; cardiavascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or Gaps /note= "phosphorothioate linkages and all cytidine ö Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:427. 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; residues are 5-methylcytidines /mod\_base= OTHER /note= "2'-O-methocyethyls" 16..20 100.0%; Pred. .... Claim 4; SEQ ID NO 198; 132pp; English. Location/Qualifiers 2709 AAAAAAAAAAAAAAAA 2728 /mod\_base= OTHER 20 AAAAAAAAAAAAAAAA 1 BP. ADM14240 standard; DNA; 20 (first entry) \*tag= a Q Local Similarity 100. /\*tag= Key modified\_base modified\_base modified base Homo sapiens 01-JUL-2004 Synthetic. ADM14240; Query Match RESULT 586 Matches ADM14240/ ð 셤 

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to human mPGES-1 gene is located on chromosome 9, more specifically to having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisabse oligonucleotides and antisense compounds have cytostatic, antidiabbetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, altherise.
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tive 0; Mismatches 0; Indels
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                    /note= "2'-0-methoxyethyls"
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to operate invention also describes: (1) antisense compounds, 9434.3. The present invention also describes: (1) antisense compounds, compGES-1, which specifically whybridise with the nucleic acid encoding compGES-1 in cells or tissues; and (3) a method of inhibiting the expression of inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal compounds a disease or condition associated with meGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisfammatory, neuroprotective, nootropic, antisfammatory, neuroprotective, nootropic, antisfammatory, neuroprotective, nootropic, antisfammatory, neuroprotective, nootropic, antistriities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or cophthalmic, immunological, cardiovascular or neurological disorder.
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tive 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "2'-0-methocyethyls"
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                                                                   ocation/Qualifiers
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ophthalmic, immunological, cardiovascular or neurological disorder.

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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SXS ð 셤 The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal mpGES-1 in cells or tissues; and (3) a method of treating an animal naving a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisidabetic, immunomodulator, cardiant, neuroprotective, ontistidabetic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Albeheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatitc; antidiabetic; immunomodulator; cardiant; neuroprotective; antidifiammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; neuroprotective; cardiant; neuroprotective; quene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se. encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or compound, having a sequence targeted to a nucleic acid /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines" Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:197. /mod\_base= OTHER /note= "2'-0-methocyethyls" 16. .20 /mod\_base= OTHER /note= "2'-0-methoxyethyls" Claim 4; SEQ ID NO 197; 132pp; English. Location/Qualifiers base= OTHER 25-SEP-2003; 2003WO-US030374 25-SEP-2002; 2002US-0413549P 01-JUL-2004 (first entry) Д \*tag= a O useful . .20 \*tag= (PHAA ) PHARMACIA CORP. \*tag≖ WPI; 2004-305094/28. WO2004028458-A2 antisense modified\_base nodified base modified\_base Homo sapiens. Synthetic. 08-APR-2004. Gierse JK; ADM14010; ischemia. New

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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic, antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antienthalmological; almunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
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                                     Gaps
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    0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
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25-SEP-2002; 2002US-0413549P.

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trargeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits lite expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidabetic, immunomodulator, cardiant, neuroprotective, antidabetic, immunomodulator, cardiant, neuroprotective, antidabetic, immunomodulator, and acation activities, and can cophthalmological, immunomodulatory and cardiavascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or dispace of a sease of the preparing a composition in the used for preparing a composition in the control of the co
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/note= "phosphorothioate linkages and all cytidine
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The present sequence represents a chimeric antisense oligonuclectide targeted to human microsomal prostaglandin E2 synthase (mFGES-1). The thuman mPGES-1 gene is located on chromosome 9, more specifically to operate the present invention also describes: (1) antisense compounds, compoused, which specifically bybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisiness oligonucleotides and antisense compounds have cytostatic, antisinhibatic, immunomodulator, cardiant, neuroprotective, copthalmological, immunomodulatory and cardiovascular activities, wasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mFGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Althebiaer's
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGBS-1). The human mpGSS-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: () antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding (mPGBS-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of treating an animal having a disease or condition associated with mPGBS-1. MPGBS-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antishabetic, immunomodulator, cardiant, neuroprotective, antihilammatory, neuroprotective, antihilammatory, neuroprotective, antihilammatory neuroprotective, antihilammatory and and a modern and a mpGBS-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or disease or disease or condition associated with MPGBS-1 e.g., inflammation, Alzheimer's
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Pred. No. 6.8e+02;
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residues are 5-methylcytidines"
                                                        /*tag= a /*tag= a //mod_base= OTHER //mod_base= OTHER 16. 20 /*tag= c //etag= c //etag= c //etag= CTHER //note= "2'-0-methoxyethyls"
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human meges-1 agene is located on chromosome 9, more specifically to of 404.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of complete in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisabetic, immunomodulator, cardiant, neuroprotective, notropic, antiarthritic, vasotropic, antinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition injury, or continued in memory of the can be used in gene chromery is cancer, isothemia or reperfusion injury, or continued in memory of the can be used in the cancer, isothemia or reperfusion injury, or continued in the cancer, isothemia or reperfusion injury, or can be used to the cancer, isothemia or reperfusion injury, or can be used to the cancer, isothemia or reperfusion injury, or can be used to the cancer, isothemia or reperfusion injury, or can be used to the cancer, isothemia or reperfusion injury, or cancer, isothemia or reperfusion injury, or can be used to the cancer, isothemia or reperfusion injury, or can be used to the cancer is the 
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Claim 4; SEQ ID NO 376; 132pp; English.
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Query Match

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0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02;
                       0; Indels
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            Local Similarity
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Gaps

2709 AAAAAAAAAAAAAAA 2728 à

chimeric, antisense oligonucleotide, phosphorothioate, human, microsomal prostaglandin E2 synthase, mPGES-1; mPGES-1 inhibitor; Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:376.

20 AAAAAAAAAAAAAAAA

ð a ADM14189 standard; DNA; 20 BP

ADM14189

(first entry)

01-JUL-2004

ADM14189;

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microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                      immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; neuroprotec
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/note= "2'-0-methocyethyls"
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       ADM13996;
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ADM13996,
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New antisense compound, having a sequence targeted to a nucleic acid
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                                                                                                                                                                                                                                            chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; implibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; noctropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:183.
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ö antidiabetic, immunomodulator, cardiant, neuroprotective, antiantial ammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder. chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss. Gaps /mod\_base= OTHER /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines" ö Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:188 Score 20; DB 1; Length 20; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; /mod\_base= OTHER /note= "2'-O-methocyethyls" 16. .20 /note= "2'-0-methoxyethyls" Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches 0 Location/Qualifiers 2709 AAAAAAAAAAAAAAAAA 2728

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The present sequence represents a chimeric antieense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human meGES-1 gene is located on chromosome 9, more specifically to ogd 4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically thybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal contribute its expression; (2) a method of tinhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal cantidiabetic, immunomodulator, cardiant, neuroprotective, antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound completed with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or condition associated with mPGES-1 e.g., inflammation in disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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.00.0%; Pred. No. 6.8e+02;
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/note= "2'-0-methoxyethyls"
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/note= "2'-0-methocyethyls"
16. .20
                                                                           Claim 4; SEQ ID NO 188; 132pp; English.
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ADM14012 standard; DNA; 20 BP

RESULT 596

ADM14012,

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunoomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunoomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.

Homo sapiens

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:199.

(first entry)

01-JUL-2004

ADM14012;

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to open solid prostaglandin E2 synthase (mPGES-1). The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically whybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal mPGES-1 in cells or tissues; and (3) a method of treating an animal artisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiamacroy, neuroprotective, nootropic, antisthammacroy, neuroprotective, nootropic, antisthammacroy, neuroprotective, nootropic, antistense compound to phthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 191; 132pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                      ischemia.
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01-JUL-2004 (first entry)
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (MPGES-1). The human mMGES-1 gene is located on chromosome 9, more specifically to operate on also describes: (1) antisense compounds, pays.3. The present invention also describes: (1) antisense compounds, naving a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of naving a disease or condition associated with mPGES-1. MPGES-1 and cardiabetic; immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiavascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alibhiator encoding a disease or condition associated with mPGES-1 e.g., inflammation, Alibhiator encoding a disease or condition associated with mPGES-1 e.g., inflammation.
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                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                         /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                        /*tag= a
/mod_base= OTHER
/note== "2'-O-methocyethyls"
                                                                                                                                                                              /note= "2'-0-methoxyethyls"
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                        Location/Qualifiers
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/mod_base= OTHER
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Best Local Similarity 100.0
wheres 20; Conservative
                                               *tag= b
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2709 AAAAAAAAAAAAAAAAA 2728

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20 AAAAAAAAAAAAAAAA 1

ADM14015 standard; DNA; 20 BP

RESULT 597 ADM14015, ADM14015

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to having a sequence compounds, 9434.3. The present invention also describes: (1) antisense compounds, paying a sequence comprising 8-30 bp targeted to a nucleic acid encoding compounds in thibits its expression; (2) a method of inhibiting the expression of inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal compounds a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, corportective, nootropic, antidialmanicy, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or composition decent as mPGES-1 inhibitors and in gene therapy. The antisense compound the used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound confition associated with mPGES-1 e-g., inflammation, Alaheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or computating, immunological, cardiovascular or neurological disorder.
                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; ophthalmological;
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                                                                                                                                                                                                                                                                        immunomodulatory, cardiovascular, gene therapy, inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia, reperfusion injury, ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:202
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/note= "2'-O-methocyethyls"
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                                           Gaps
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                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:208
                   Query Match 0.7%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0; Indels
Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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/note= "2'-0-methocyethyls"
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/note= "2'-0-methoxyethyls"
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                       having a sequence comprising 8.30 bp targeted to a nucleic acid encoding mPCES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and cardiovascular activities and can ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound condition associated with mPGES-1 e.g., inflammation, alternating a disease or condition associated with mPGES-1 e.g., inflammation, alternating disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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The present invention also describes: (1) antisense compounds
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/note= "2'-0-methocyethyls"
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Matches 20; Conservative
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Gierse JK;
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                                                                                                                                                                                      The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human megES-1 gene is located on chromosome 9, more specifically to open is located on chromosome 9, more specifically to open is located on chromosome 9, more specifically to open in the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal condition associated with mpGES-1. MpGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antidiamaticity, namenomodulator, and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound to be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used as used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alabeimer's condition associated with mpGES-1 e.g., inflammation, Alabeimer's condition associated with mpGES-1 e.g., inflammation, and injury, or oppthhalmic, immunological, cardiovascular or neurological disorder.
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                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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Pred. No. 6.8e+02;
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0.7%; Score 2v; C.

Best Local Similarity 100.0%; Pred. No. 6.8

Matches 20; Conservative 0; Mismatches
                                                                                                                                                                 Claim 4; SEQ ID NO 575; 132pp; English.
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   (PHAA ) PHARMACIA CORP.
                                                          WPI; 2004-305094/28
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                                Gierse JK;
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to pag4.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal companies oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisinglammatory, neuroprotective, nootropic, antianflammatory, neuroprotective, nootropic, antianflammatory, neuroprotective, nootropic, antianflammatory, neuroprotective, and candidate as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Althebiaer's
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
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/*tag= a
/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
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/note= "2'-O-methoxyethyls"
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Best Local Similarity
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                                                                                                                 modified base
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chimeric; antisense oligonuclectide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGBS-1; mPGBS-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antidiformaneory; neuroprotective; antidiformaneory; neuroprotective; neuroprotective; osotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;

Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:274.

(first entry)

01-JUL-2004

ADM14087;

ADM14087 standard; DNA; 20 BP.

RESULT 602

/note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"

Location/Qualifiers

/mod\_base= OTHER

Ω

\*tag=

modified\_base

Homo sapiens.

Synthetic.

/note= "2'-0-methoxyethyls"

WO2004028458-A2.

38-APR-2004.

25-SEP-2003; 2003WO-US030374. 25-SEP-2002; 2002US-0413549P.

(PHAA ) PHARMACIA CORP.

base= OTHER

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modified base

/note= "2'-0-methocyethyls'

/mod base= OTHER

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is a located on chromosome 9, more specifically to off an expecifically the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the mucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal antisense oligonucleotides and antisense compounds have cytostatic, antidiabbetic, immunomodulator, cardiant, neuroprotective, antidiabbetic, immunomodulatory and cardiant, neuroprotective, antidiabbetic, immunomodulatory and cardiant activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, sand, and sand and sand antisense of condition associated with mpGES-1 e.g., inflammation, sand, san
neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetees; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
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ophthalmic, immunological, cardiovascular or neurological disorder.
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16. .20
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/mod_base= OTHER
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGBS-1). The human mPGBS-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGBS-1, which specifically hybridise with the nucleic acid mPGBS-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGBS-1 in cells or tissues; and (3) a method of treating an animal atsisense or condition associated with mPGES-1. MPGBS-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic,
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Gaps

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Length 20; 0; Indels

Score 20; DB 1; Le Pred. No. 6.8e+02;

0.75, 100.0%; Pr.

Conservative

Query Match Best Local Similarity Matches 20; Conserv

Mismatches

2709 AAAAAAAAAAAAAAAA 2728

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                compound
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                                                                disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, ophthalmic, immunological, cardiovascular or neurological disorder.
ophthalmological, immunomodulatory and cardiovascular activities, and be used as mFGES-1 inhibitors and in gene therapy. The antiseense component be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                            Gaps
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                                                                                                                                            DB 1; Length 20; 6.8e+02;
                                                                                                                                                                           0; Indels
                                                                                                            Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "2'-0-methocyethyls"
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                                                                                                                                          Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (meGES-1). The human meGES-1 special socated on chromosome 9, more specifically to human meGES-1 special socated on chromosome 9, more specifically to antisense compounds, 944.3. The present invention also describes: (1) antisense compounds, compGES-1, which specifically hybridise with the nucleic acid meGES-1 and inhibits its expression; (2) a method of inhibiting the expression of meGES-1 in cells or tissues; and (3) a method of treating an animal compounds a disease or condition associated with meGES-1. MeGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, contininamatory, neuroprotective, nootropic, antistriatic, vasotropic, antistrialmamological, immunomodulatory and cardiovascular activities, and can be used as meGES-1 inhibitors and in gene therapy. The antisense compound can be used as meGES-1 inhibitors and in gene therapy. The antisense compound con be used as meGES-1 inhibitors and in gene therapy. The antisense compound con be used as meGES-1 inhibitors and in gene therapy. Alzheimer's condition associated with meGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or condition associated with meGES-1 e.g., inflammation injury, or condition immunological, cardiovascular or neurological disorder.
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Pred. No. 6.8e+02;
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/note= "2'-O-methocyethyls"
16. .20
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                                  Claim 4; SEQ ID NO 487; 132pp; English.
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/mod_base= OTHER
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Matches 20; Conserv
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modified_base
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-10 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibite its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal national adjouncleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, optibalmological, immunomodulatory neuroprotective, notropic, antidiamatory, neuroprotective, notropic, antidiamatory interoperating a composition be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer, isohaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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100.0%; Pred. No. 6.8e+02;
ve 0; Mismatches 0; Indels
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                                                             25-SEP-2003; 2003WO-US030374.
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Synthetic.
                      08-APR-2004,
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                                                                                                                                                                                                                                                                                                                                                      ischemia.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to offer an engine of antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding compense. Which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal companies of antisense oligonucleotides and antisense compounds have cytostatic, antishabetic, immunomodulator, cardiant, neuroprotective, antishabetic, immunomodulatory and cardiant, neuroprotective, antishabetic, immunomodulatory and cardiant, neuroprotective, cophthalmological, immunomodulatory and cardiavascular activities, and can be used for preparing a omposition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Altheimer's conditions.
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                                                                              /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
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100.0%; Pred. No. 6.8e+02;
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/note= "2'-O-methoxyethyls"
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/note= "2'-O-methocyethyls'
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Location/Qualifiers
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Key
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; neuroprodulator; cardiant; neuroprotective; antidilamanory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzaelmer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; per cardiovascular disorder; neurological disorder; se. New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or /mod\_base= OTHER /note= "phosphorothioate linkages and all cytidine Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:194 chimeric; antisense oligonucleotide; phosphorothioate; human; residues are 5-methylcytidines /mod\_base= OTHER /note= "2'-O-methocyethyls" 16. .20 /note= "2'-0-methoxyethyls' Location/Qualifiers mod\_base= OTHER 25-SEP-2003; 2003WO-US030374 25-SEP-2002; 2002US-0413549P ..20 \*tag= b \*tag≃ a \*tag= c (PHAA ) PHARMACIA CORP WPI; 2004-305094/28. WO2004028458-A2 Key modified\_base modified\_base modified\_base Homo sapiens. Synthetic. 08-APR-2004 Gierse JK;

schemia.

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Claim 4; SEQ ID NO 194; 132pp; English.

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The present sequence represents a chimeric antisense oligonucleotide
targeted to human microsomal prostaglandin E2 synthase (mCGES-1). The
thuman mPGES-1 gene is located on chromosome 9, more specifically to
the present invention also describes: (1) antisense compounds,
companies, which specifically hybridise with the nucleic acid encoding
mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and
inhibits its expression; (2) a method of inhibiting the expression of
inhibits its expression; (2) a method of inhibiting the expression of
inhibits its expression; (2) a method of inhibiting the expression of
inhibits its expression; (2) a method of inhibiting the expression of
antisense oligonucleotides and antisense compounds have cytostatic,
antisense oligonucleotides and antisense compounds have cytostatic,
antisflammatory, neuroprotective, nootropic, antiathritic, vasotropic,
antisflammatory, neuroprotective, nootropic, antiathritic, vasotropic,
ophthalmological, immunomodulatory and cardiovascular activities, and can
be used as mPGES-1 inhibitors and in gene therapy. The antisense compound
can be used for preparing a composition for treating a disease or
condition associated with mPGES-1 e.g., inflammation, Alzheimer's
disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
cophthalmic, immunological, cardiovascular or neurological disorder.

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; eyrostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9d34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                               ;;
0
                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:311.
DB 1; Length 20; 6.8e+02;
                               0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mod_base= OTHER
/note= "2'-0-methocyethyls"
   0.7%; Score 20; DB 100.0%; Pred. No. 6.8 :ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2'-0-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                              2709 AAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     base= OTHER
                                                                                         20 AAAAAAAAAAAAAAAA 1
                                                                                                                                                                 BP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-SEP-2003; 2003WO-US030374
                                                                                                                                                                 ADM14124 Standard; DNA; 20
                                                                                                                                                                                                                         (first entry)
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                               Conservative
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      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified base
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                                                                                                                                                                                                                         01-JUL-2004
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                                 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                               ADM14124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ischemia.
                   Best Loca
Matches
                                                                                                                                     RESULT 607
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                                                                                                                                                    ADM14124,
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mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidialentic, immunomdulator, cardiant, neuroprotective, ophthalmological, immunomdulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimeric; antisense oligonuclectide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inplibitor; cytocatatic; antidiabetic; immunoomodulator; cardiant; neuroprotective; antidinflammatory; neuroprotective; autidinflammatory; neuroprotective; vasotropic; ophthalmological; inmunoomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                  Gaps
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:403
                                                                                                                                                                                                                                                                                            Query Match 0.7%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 6.8e+02; Atches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
/note= "2'-0-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                      20 AAAAAAAAAAAAAAAAA 1
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modified_base
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                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 608
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mepgES-1 gene is located on chromosome 9, more specifically to office the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antialabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antinflammatory, neuroprotective, nootropic, antiantises, and can can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Raman label; specific binding member; surface-enhanced Raman scattering;
                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0%; Pred. No. ....
Marches 20; Conservative 0; Mismatches
                                                                                                                                                        Claim 4; SEQ ID NO 403; 132pp; English.
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28-MAY-2002; 2002US-0383630P.
14-JUN-2002; 2002US-00172428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAY-2003; 2003US-00431341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                  WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-418413/39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2004086897-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CAOY/) CAO Y.
(JINR/) JIN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004.
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 Gierse JK;
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                                                                                                                         ischemia.
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WPI; 2004-440357/41.

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Mucic RC, Storhoff JJ, Elghanian R;
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Garimella V, Li Z;
                                                                                                                                                                                                                                                                 ADP20152 standard; DNA; 20 BP
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29-JAN-1999; 99US-00240755.
25-UNN-1999; 99US-00344667.
13-JAN-2000; 2000US-0176409P.
                                                                                                                                                                                                                                                                                                                                                                                                                                     26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
26-JUN-2000; 2000US-0213906P.
                                                                                                                                                                                                                                                                                                                                                                                        18-NOV-2003; 2003US-00716829.
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                                                                                                                                                                                                                                                                                           26-AUG-2004 (first entry)
                                                                                                                                                                                                     Conservative
                          specific binding member
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NANO-) NANOSPHERE INC
                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                             US2004110220-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Taton TA,
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The invention describes a reagent (I) comprising a particle bound to at least one Raman label and a specific binding member, where the Raman cartivated to provide a surface-enhanced Raman scattering (SERS) effect or comprising a specific binding member having two or more different Raman labels bound to it. Also described are: a test kit (II), comprising (I) in one container and a silver, gold or copper Raman container and a silver, gold or copper Raman compares that in another container, and a fibre optic detection device (III), having a bundle of optical fibres terminating with ends of the optical fibre, where a several of the optical fibres have (I) located at the ends of the optical fibre. (I) is useful for: detecting for the presence or absence of one or more target analytes in a sample, the carget analytes having at least two binding sites; detecting the presence or absence of one or more target analytes in a sample, the sequence or absence acid aving at least two binding sites; detecting the presence or absence acid aving at least two portions; and for sereening one or more molecules to determine whether the molecule is a ligand to one or more specific receptors. This sequence represents an oligonucleotide associated with the SERS-based detection analyte detection method.
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Reagent, useful for detecting target analyte e.g., nucleic acid, comprising particle having bound to at least one Raman label, which can be activated to provide surface-enhanced Raman scattering effect, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 31; 55pp; English.
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Line inventions to tailly a considered of detecting a nucleic acid with at conjugate, contacting the nucleic acid and nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow the providisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the hybridisation. The aligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. The method is used for detection and separation of nucleic acids (e.g. viral DNA, a gene associated with a disease.)

Cf nucleic acids (e.g. viral DNA, a gene associated with a disease.)

Cf nucleic acids (e.g. viral DNA, synthetic DNA, structurally-modified DNA, DNA from biological sources or PCR products) for diagnosis of various diseases (such as genetic diseases, bacterial infections and viral confections) and for forensics, DNA sequencing, paternity testing and monitoring gene therapy. This sequence represents a linking and conformation.
                                                    articles useful for detection and separation of nucleic acids e.g associated with disease, in a diagnostic assay, comprise several
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gapa
                                                                                                                                                                             The invention relates to a method of detecting a nucleic acid with
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                       Example 24; SEQ ID NO 70; 142pp; English.
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                                                                                           oligonucleotides attached to them.
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Taton TA, Garimella V, Li 2;
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13-JAN-2000; 2000US-0176409P,
26-APR-2000; 2000US-0200161P,
26-JUN-2000; 2000US-00603830.
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99US-00240755
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2000US-0213906P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.º
Matches 20; Conservative
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                                                         Nanoparticles useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JUL-1997;
29-JAN-1999;
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89US-00422383.
90US-00537198.
90US-00573616.
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                                                                                                                                                                                        16-OCT-1989;
                                                                                                                                                                                          11-JUN-1990;
24-AUG-1990;
                                                                                                                                                                             06-JUL-2004
                                                                                                           ADP99303;
                                                                         Query Match
                                                                                                                                                                  Mammalia.
                                                                              Matches
                                                                                                 RESULT 612
                                                                                                    ADP99303,
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conjugate, contacting the nucleic acid and nanoparticle-oligonuclectide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonuclectides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonuclectides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonuclectides on the nanoparticles with the nucleic acid results in a detectable change. The method is used for detection and separation of the oligonuclectides on the method is used for detection and separation. Of nucleic acids (e.g. viral DNA, a gene associated with a disease, bacterial DNA, fungal DNA, synthetic DNA, structurally-modified DNA, DNA from biological sources or FCR products) for diagnosis of various diseases (such as genetic diseases, bacterial infections and viral monitoring gene therapy. This sequence represents a linking and monitoring diseases.
                                                                                                              Nanoparticles useful for detection and separation of nucleic acids e.g. genes associated with disease, in a diagnostic assay, comprise several oligonucleotides attached to them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of detecting a nucleic acid with at
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                                                                                                                                                                                                                                                                                                                                                 Example 18; SEQ ID NO 55; 142pp; English.
WPI; 2004-440357/41.
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invention relates to preparing a (vertebrate) human stem cell factor

hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.

Example 3; SEQ ID NO 33; 210pp; English.

Preparing a human stem cell factor (SCF) polypeptide, useful for

Suggs SV, Martin FH;

Bosselman RA,

Zsebo KM,

(AMGE-) AMGEN INC.

WPI; 2004-497128/47.

90US-00589701. 91US-00684535. 92US-00982255. 93US-00172329. 95US-00449182.

10-APR-1991; 25-NOV-1992; 21-DEC-1993; 01-OCT-1990;

24-MAY-1995;

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Gaps
                                     ;
0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
rative 0; Mismatches 0; Indels
                                                                   2709 AAAAAAAAAAAAAAAA 2728
                                                                                                    1 AAAAAAAAAAAAAAA 20
                Local Similaricy
nes 20; Conservative
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SCF; stem cell factor; gene therapy; haematopoietic progenitor cell; aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Letterer-Siwe disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  congestive splenomegaly, Kala awar; sarcoidosis; primary splenic pancytopaenia; milary tuberculosis, diseminated fungus disease; Fulminating septicaemia; malaria; vitamin B12 deficiency; folic acid deficiency; pyridoxine deficiency; biamond Blackfan anaemia; hypopigmenteation disorder; piebaldism; vitiligo; neurological damage; infertility; intestinal damage; irradiation; chemotherapy; AIDS; haematopoletic recovery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  refractory erythroblastic anaemia; Di Guglielmo syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acute blood loss; neoplasm; cancer; ss; PCR; primer
                                                                                                                                                                                                                                                                                                        Stem cell factor, SCF, universal PCR primer #3.
ADP99303 standard; DNA; 20 BP
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The invention fatares to preparing a (vertexate) man size metil factor (SCP) polypeptide comprising growing host cells transformed or transfected with DNA encoding a human SCP that stimulates growth of heematopoietic progenitor cells unrised included is a recombinant host cell polypeptide produced. Also included is a recombinant host cell caracter of transfermed or transferred with an expression control sequence, and isolating the operatively linked to a mexpression control sequence, and isolating the cransferred of transferred of the vertebrate SCP polypeptide encoding DNA operatively linked to a member of the vertebrate SCF polypeptide in the host cell. Disclosed as new are rated the vertebrate SCF polypeptide in the host cell. Disclosed as new are rated to the vertebrate SCF polypeptide in the host cell. Disclosed as new are rated to recombinantly expressed SCF protein fragments. The expression of sequences are useful for effecting the large scale synthesis of SCF by a current of sequences are useful for effecting the large scale synthesis of SCF by a current of sequences are useful for effecting the large scale synthesis of SCF by a current of such host cells, and new and useful viral and circular plasmid DNA vectors, new and useful as and circular plasmid DNA vectors. The DNA sequences are also useful as and its related probes; in soletting human spendic DNA encoding SCF, in methods for production of SCF and SCF products in quantity. The SCF is soletting for production of SCF and SCF products in quantity. The SCF is useful for treating heematopoletic disorders, e.g., aplastic anaemia, consistent of story productions and sequences are used. The scription of sequences are used. The scription of sequences are used. The scription of sequences of productions, and the sequences are subscriptions, and sequences are subscriptions. The scriptions are subscriptions, and sequences are also useful for treating neurological damage, infertility states, intentinal damage for each inmune system for tighting neoplasia, cau
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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Best Local S
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myelosuppression; immune deficiency; mecplasm; nerve damage; infertility; intestinal damage; myeloproliferative disorder; early haematopoietic progenitor cell; haematopoietic disorders; applastic ansemia; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Diamond-Blackfan anaemia; DBA; Fanconi's anaemia; gene therapy; acute blood loss; 88; PCR; primer;

89US-00422383. 90US-00537198. 90US-00533616. 91US-00689701. 91US-00684535. 92US-00982255.

10-APR-1991; 25-NOV-1992; 21-DEC-1993; 07-JUN-1995;

95US-00486546.

19-JUN-2002; 2002US-00175608

16-OCT-1989;

11-JUN-1990 24-AUG-1990; 01-OCT-1990;

Rattus norvegicus.

sapiens

probe Ношо US2004181044-A1.

16-SEP-2004

anaemia; bone marrow during transplant; bone marrow aplasia;

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The invention relates to a novel method for isolating molecules using a micro-channel. The molecules are isolated by introducing a mixed solution having two types of solute molecules differing in molecular weight into a micro flow path, to form a non turbulent flow, and providing physical action to the molecules by changing the flow state, thus causing different behaviours among different solute molecules, where the different behaviour enables uneven distribution of specific kinds of molecules in the flow path, causing separation of specific kinds of invention further comprises molecule separation of specific kinds of unbettact with a micro flow path, having one or more curved portions, a sample intake unit at one side and a sample removal opening at the other side, and a physical property detection sensor arranged inside the curved portion. The method is useful for isolating molecules, e.g. polymer compounds, DNA or proteins. The method enables might and efficient separation of molecules by utilising specific flow behaviour in a non turbulent flow, in a micro flow path, where a large number of samples can be processed. This polymucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                Isolating molecules e.g., DNA, by introducing solution with two types of solute molecules into micro flow path to form non turbulent flow, providing physical action to molecule causing difference in flow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stem cell factor; SCF; haematopoietic; HT1080 fibrosarcoma cell line;
5637 bladder carcinoma cell line; leukopaenia; thrombocytopaenia;
                                                                    molecule isolation; micro-channel; molecular weight; micro flow path; polymer compound; flow behaviour; non turbulent flow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                         Yamashita K, Maeda H, Shimizu H, Miyazaki M, Nakamura H;
Yamaguchi Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
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                                      Micro-channel molecule isolation related Adenine oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                           (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 7; 19pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           behavior, separating molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADU50633 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%;
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                                                                                                                                                                                                                                                                          18-FEB-2003; 2003JP-00039870.
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hes 20; Conservative
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                                                                                                                               Unidentified.
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The invention relates to a stem cell factor (SCF) such as non-naturally-
ccurring SCF having an amino acid sequence sufficiently duplicative of
that of naturally occurring SCF to allow possession of a haemacopoletic
biological activity of naturally occurring stem cell factor, or naturally
ccurring SCF. Also included are an isolated DNA sequence for use in
securing SCF. Also included are an isolated DNA sequence for use in
ccurring expression in a prokaryotic or eukaryotic host cell of non-
cc naturally occurring SCF, a prokaryotic or eukaryotic host cell.
cransformed or transfected with the DNA, a polypeptide product of the
expression of the DNA in a prokaryotic or eukaryotic host
cc expression of the DNA in a prokaryotic or eukaryotic host
cc expression of non-naturally occurring SCF, a DNA sequence coding for a
polypeptide fragment or polypeptide analogue of naturally-occurring stem
cc expression of non-naturally occurring SCF, a DNA sequence coding for a
polypeptide fragment or polypeptide analogue of naturally-occurring stem
cc expression of non-naturally occurring SCF, a DNA sequence coding for a
ccll factor, a biologically functional plasmid or viral DNA vector
ccll factor, a biologically the DNA, a polypeptide having part
cc carcinoma cell line, or human SCF cDNA obtained from HT1080
cc fibrosarcoma cell line, or human SCF cDNA obtained from HT1080
cc fibrosarcoma cell line, or human SCF cDNA obtained from HT1080
cc fibrosarcoma cell line, or human SCF cDNA obtained from HT1080
cc fibrosarcoma cell line, or human SCF cDNA obtained from Gradiact
cc carcinoma cell line, or mammal. SCF is useful for treating patients
cc carcinoma cell line, and enhancing engraftment of bone marrow
during transplantation in a mammal. SCF is useful for treating patients
cc recovery in treatment of radiation, chemical, or chemotherapeutic in paramally effective doses of SCF. SCF is useful for treating
cc with therapeutically effective doses of SCF. SCF is useful for treating
cc intestinal damage, and a myeloproliferative disor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel stem cell factor (SCF) such as non-naturally-occurring SCF or naturally occurring SCF, useful for treating leukopenia, thrombocytopenia, anemia, and enhancing engraftment of bone marrow during
                                                                                                                                                                                                                                                                                                                                                                        Martin FH;
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07-AUG-2000; 2000US-00635249.
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                                                                                                                                                        BOSSELMAN R A.
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(MART/) MARTIN F H.
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                                                                                                             ZSEBO K M.
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(BOSS/) E
(SUGG/) S
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transfecting early haematopoietic progenitor cells with a gene which involves culturing early haematopoietic progenitor cells with SCF, and transfecting the cultured cells with a gene. SCF is useful for transfecting the cultured cells with SCF, transfecting the cultured cells with a gene, and administering the cultured cell to the mammal. SCF is useful for treating various haematopoietic disorders, aplastic anaemia, myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute disease, Niemann-Pick disease, Diamond-Blackfan anaemia (DBA), Fanconi's anaemia. SCF is useful for enhancing the efficiency of gene therapy, for enhancing haematopoietic recovery after acute blood loss. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid detection; genetic disease; cystic fibrosis; buchenne muscular dystrophy; sickle cell anaemia; phenylketonuria; bacterial disease; tuberculosis; Lyme disease; viral disease; microbial infection; sexually transmitted disease; gonorrhoea; forensics; DNA sequencing; paternity testing; cell line authentication;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of nucleic acid by contacting nucleic acid with substrate including oligonucleotides, contacting nucleic acid bound to substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thiol-modified oligo (SA20) to form oligo-nanoparticle conjugates.
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0
                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                           0; Indels
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                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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Pred. No. 6.8e+02;
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100.0%; Pred. No. ...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                     2707 CTAAAAAAAAAAAAAA 2726
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/mod_base= OTHER
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97WO-US012783.
99US-00240755.
99US-00344667.
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                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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21-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADU17674;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taton TA;
                                                                                                                                                                                                                                                                                                                                                                                               RESULT 615
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Gaps

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The present invention provides methods of detecting nucleic acids. The method of the invention comprises contacting a nucleic acid with a comparises contacting a nucleic acid with a compariate including oligonucleotides under conditions to allow hybridisation of oligonucleotides on the substrate, contacting nucleic acid bound to the substrate with nanoparticles comprising of conductivity. The invention is useful for detecting nucleic acids for the conductivity. The invention is useful for detecting nucleic acids for the diagnosis of genetic diseases such as cystic fibrosis, buchenne muscular conductivity. The invention is useful for detecting nucleic acids for the diagnosis of genetic diseases such as cystic fibrosis, buchenne muscular cuch stuberculosis, Lyme disease, Helicobacter pylori infections.

CE Scherichia coli infections, viral diseases such as human immunodeficiency and Salmonella infections, viral diseases such as human immunodeficiency disease virus, (RIV), heptitis virus, herpes virus, cytomegalovirus and Epstein-Barr virus, sexually transmitted disease such as genorrhoea The invention is also useful in forensics, DNA sequencing, paternity testing, cell line authentication and gene therapy. The present sequence is a thiol-modified coligonucleotide is used to detect the nanoparticle conjugates. This oligonucleotide is used to detect the effect of coadsorbed diluent oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
nanoparticles comprising oligonucleotides, and detecting change in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.7%; Score 20; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Allergic response suppressor oligonucleotide #560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Prec. ...
                                                    Example 18; SEQ ID NO 55; 110pp; English.
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2001US-00776479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADU89876 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20; Conservative
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PETERSEN D M
FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                     conductivity.
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02-FEB-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-NOV-2004.
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(FOUR/)
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nucleic acid alone or in combination with other

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Gaps

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Query Match
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medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; 20; Conservative method of the invention. an immunostimulatory Query Match Best Local Similarity Synthetic. ADU89542; RESULT 618 Matches ADU89542/ 8888888 ò 셤 ö The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and atopic dermatitis, using The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and atopic dematitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second do Gaps 98; antiasthmatic; antiallergic; dermatological; antinflammatory; antibacterial; virucide; immunoglobulin E antagonier; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection. ö 0.7%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 6.8e+02; 0; Indels Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; Allergic response suppressor oligonucleotide #556. 0; Mismatches Disclosure; SEQ ID NO 556; 235pp; English. Disclosure; SEQ ID NO 560; 235pp; English dermatitis, in a subject, comprises a of an immunostimulatory nucleic acid. Fouron Y; 2709 AAAAAAAAAAAAAAAA 2728 АААААААААААААААА 20 BP. 100.08; 03-FEB-2000; 2000US-0179991P. 23-APR-2004; 2004US-00831778 ADU89872 standard; DNA; 20 Petersen DM, (first entry) Matches 20; Conservative nethod of the invention. 'nΣ WPI; 2004-833006/82. (BRAT/) BRATZLER R (PETE/) PETERSEN D Best Local Similarity (FOUR/) FOURON Y. US2004235774-A1 Bratzler RL, 10-FEB-2005 25-NOV-2004 Synthetic. ADU89872;

ö The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allery, including rhinitis, uriticaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid. Gaps ss; antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection. ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; iive 0; Mismatches 0; Indels Seguence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other; Allergic response suppressor oligonucleotide #226. Disclosure, SEQ ID NO 226; 235pp; English. Fouron 2709 AAAAAAAAAAAAAAAAA 2728 20 AAAAAAAAAAAAAAAAAA 1 ВЪ. 03-FEB-2000; 2000US-0179991P. 02-FEB-2001; 2001US-00776479. 23-APR-2004; 2004US-00831778 Ğ, ADU89542 Btandard; DNA; 20 (first entry) Query Match 0.7 Best Local Similarity 100. Matches 20; Conservative method of the invention. Petersen (BETE/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y. WPI; 2004-833006/82. US2004235774-A1. 10-FEB-2005 Bratzler RL, 25-NOV-2004. 

2709 AAAAAAAAAAAAAAAAA 2728

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respiratory syndrome (SARS) coronavirus. The polypeptides include spike (S or E2), env (E or SM), membrane (M or E1), hemagglutinin-esterase (HE or E3), and nucleocapsid (N) polypeptides, and the ORFiab and ORFiab (replicase) polypeptides and their proteolytic fragments. The invention also relates to antibodies which recognise the polypeptides; nucleic acids encoding the SARS virus polypeptides; primers specific for SARS virus nucleic acids adouble-stranded RNA molecule 10-30 nucleotides in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      οĘ
                                                                                                                                                                          Vaccine, nucleic acid vaccine, drug screening, diagnosis,
SARS coronavirus infection, infection, respiratory disease, virucide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated polypeptide e.g. spike polypeptide, Env polypeptide, o
severe acute respiratory syndrome virus (SARS), useful as vaccine for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Han J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated polypeptides of the severe acute
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Donnelly JJ;
                                                                                                                                                 (dT)20 reverse transcription primer, SEQ:7389.
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Weiner A, Houghton M, Song HC, Seo MY,
Valiante N;
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2003US-0463668P.
2003US-0463983P.
2003US-0463971P.
2003US-0464838P.
                                                                       ACL79852 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                          2003US-0464899P.
2003US-0465273P.
2003US-0465535P.
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                                                                                                                        16-JUN-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR ) CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-766863/75.
                                                                                                                                                                                                                                                    WO2004092360-A2.
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23-APR-2003;
24-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                         16-APR-2003;
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22-APR-2003;
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22-MAY-2003;
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                                                                                                                                                                                                                                                                                                                                                                                14-APR-2003;
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Klenk HD,
                                                                                                                                                                                                    primer; ss.
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                                                                                                                                                                                                                            Synthetic.
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                                                                                                ACL79852;
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cc which is able to inactivate the SARS virus in a mammalian cell; an expression construct for recombinant expression of a SARS virus spike protein; a viral vector for in vivo delivery of a SARS virus polypeptide-cc encoding nucleic acid; and a mammalian cell line stably expressing a SARS viral antigen. The invention additionally provides a vaccine for the treatment or prevention of SARS comprising an inactivated SARS virus, a crimated SARS virus, a crimated SARS virus, a split SARS virus and pha-virus or preparation, or at least one purified SARS virus antigens; methods of preparation, or at least one purified SARS virus antigens; methods of comprising one or more SARS virus antigens; and a vaccine comprising one or more SARS virus antigens and one or more SARS virus antigens and one or more satisfying a therapeutically active agent by measuring the effect of the comprising one or more satis virus polypeptides and concluded acids can be used in the preparation and manufacture of vaccines for the treatment or prevention of SARS virus spolypeptides, antibodies against them, and SARS virus specific primers and kits containing them are useful for diagnosing or identifying the presence of SARS in a biological sample. The present sequence represente a primer for SARS in a biological sample. The present sequence represente a primer for this patent did not form part of the printed specification, but was containing the interparation of SARS genomic RNA. Note: The sequence data
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; hybridization; diagnosis; genetic disorder; bacterial infection; viral infection.
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Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide of the invention SEQ ID NO:55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. nc.
                                                                                                                                                                                                                                                                                                                                                                                                                       ftp.wipo.int/pub/published_pct_sequences
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99US-00240755.
99US-00344667.
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26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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29-JAN-1999;
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Nanoparticle for detecting nucleic acids for diagnosis of genetic, bacterial, and viral diseases, comprises oligonucleotides containing at

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Elghanian

Storhoff JJ,

Mucic RC,

Mirkin CA, Letsinger RL,

Faton TA;

WPI; 2005-019280/02.

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The invention relates to a novel nanoparticle having oligonucleotides attached thereto, comprising at least one type of recognition oligonucleotides, each having a spacer portion bound to the nanoparticle and a recognition portion having a sequence complementary to portion(s) of sequence of nucleic acid or another oligonucleotide. In the presence conditions, the nanoparticle forms a complex with the nucleic acid or another oligonucleotide and under hybridization conditions, the nanoparticle forms a complex with the nucleic acid or another oligonucleotide. The complex has a sharp melting profile and an increased melting temperature, relative to a melting profile and an increased melting temperature, relative to a melting profile and a not another oligonucleotide and an unlabeled or fluorophore-labeled or insertions, deletions, and/or mismatches in the nucleic acid or another oligonucleotide with a sequence identical to the oligonucleotides bound to the nanoparticle. The nanoparticles of the invention are useful for detecting nucleic acids for diagnosis of genetic, bacterial, and viral diseases. The nanoparticle is stable at elevated temperature and high salt concentration. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel glycosyltransferase capable of transferring N-acetyl-D-galactosamine (GalNAc) residue to GalNAc receptor substrate from GalNAc donor substrate, useful in screening substances that promotes/inhibits glycosyltransferase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     glycosyltransferase; N-acetyl-D-galactosamine; GalNAc; screening; ss;
least one type of recognition oligonucleotides, each having spacer portion and recognition portion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human glycosyltransferase pENTR/DTOPO vector 5' primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (DOKU-) DOKURITSU GYOSEI HOJIN SANGYO GIJUTSU SO.
(SEGK ) SEIKAGAKU KOGYO CO LID.
                                                            Example 18; SEQ ID NO 55; 108pp; English
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Best Local Similarity 100.
Matches 20; Conservative
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The invention relates to a novel glycosyltransferase capable of transferring an N-acetyl-D-galactosamine (GalNAc) residue to a GalNAc creeptor substrate from a GalNAc donor substrate: The glycosyltransferase comprises a polypeptide having sequence AU94808 containing amino acids are in the specification, or AD94808 in which one or more amino acids are substituted, deleted, inserted or rearranged. The invention further comprises: a nucleic acid encoding the 601 amino acid glycosyltransferase comprises: a nucleic acid encoding the 601 amino acid glycosyltransferase comprises: a nucleic acid sequence AD94807 having bases 12-1806 or 1-1806 of a fully defined sequence AD94807 having bases 12-1806 or 1-1806 of a fully defined sequence Complementary to AD94807; a nucleic acid capable of hybridizing under stringent conditions, with the nucleic acid capable of hybridizing under stringent conditions, with the nucleic acid capable of hybridizing the glycosyltransferase encoding DNA or its complementary sequence complementary to the 1806 bp continuody capable of specifically recognizing the glycosyltransferase protein; an active regulator of the glycosyltransferase protein; an active regulator of the glycosyltransferase protein; an active regulator of the glycosyltransferase protein as an active regulator of the glycosyltransferase protein as an active ingredient. The glycosyltransferase complementary plan is useful as a probe for detecting in vivo expression of the glycosyltransferase. The glycosyltransferase complementary of glycosyltransferase complementary plan is useful as a probe for detecting in vivo expression of the glycosyltransferase complementary is useful as a cradent or the glycosyltransferase protein, is useful as a cradent or the glycosyltransferase protein, is useful as cative regulator of the glycosyltransferase protein, is useful as cative regulator of the glycosyltransferase protein, is useful as cative regulator of the glycosyltransferase protein, is useful as cative regulator of the glycosyltransferase ö glycosyltransferase protein. The glycosyltransferase protein is capable of transferring GalnAc residue to a GalnAc receptor substrate from a GalnAc donor substrate. This polynucleotide sequence represents a primer used in the exemplification of the invention. Gaps ö 0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels /note= "5'-steroid disulfide linker" Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other; Gene expression; DNA detection; probe; ss. Target RNA detecting detection probe. Location/Qualifiers 2709 AAAAAAAAAAAAAAAA 2728 /\*tag= a /mod\_base= other 1 AAAAAAAAAAAAAA 20 BP. 27-FEB-2003; 2003US-0450268P. 27-FEB-2004; 2004WO-US006273 ADW02146 standard; DNA; 20 (first entry) Local Similarity 100. Les 20; Conservative (NANO-) NANOSPHERE INC. WO2005001143-A2. Key modified\_base 24-MAR-2005 06-JAN-2005 Synthetic. ADW02146; Query Match RESULT 622 Matches ADW02146 ò g ö

27-FEB-2003; 2003US-0450268P.

(NANO-) NANOSPHERE INC. Mueller UR;

Bao YP,

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The invention relates to detecting or quantifying gene expression in a sample having unlabeled target nucleic acids. The method involves can be abstrate having types of capture nucleic acid sequences attached to it in an array for the detection of multiple portions of a target nucleic acid, the detection of multiple different target nucleic acids, or both, providing namoparticles having a sequence of it, the oligonucleotides bound to the namoparticles having a sequence that is complementary to at least a portion of the oligonucleotide tail; contacting the sample, the substrate, and the namoparticles, the contacting under conditions for hybridization of the target nucleic acids to the capture nucleic acid sequences bound to the substrate and hybridization of the target nucleic acids to the namoparticles, and observing a detectable change. The target cDNAs, namoparticles, and observing a detectable change. The target cDNAs with the oligonucleotides bound to the substrate are contacted simultaneously under conditions conditions councied acid sequences comprises and with the capture nucleic acid sequences comprises and oligonucleotide, cDNA, or genomic sequence fragment. The method is useful for detecting/quantifying global sequence fragment. The method is useful for detecting/quantifying global captaneously the present sequence represent abeling and target captare amplification. The present sequence represents a detection which in a sample of unlabeled target nucleic acids acids acids associated with fluorescent labeling and target and another sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification. The present sequence represents a detection probe that can be used to detect target RNA sequences. The probe comprise a steroid disulfide linker at the 5'-end followed by a recognition sequence.
                                                                            Detecting/quantifying gene expression in a sample of unlabeled target nucleic acids by contacting the sample, a substrate having capture nucleic acid sequences and nanoparticles having bound oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                 Example 1; Page 28; 54pp; English.
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/mod base= other
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Best Local Similarity
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The 20; Conservative
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  Mueller UR;
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                                                                                                                                           for hybridization.
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modified_base
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  Bao YP,
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The invention relates to detecting or quantifying gene expression in a sample having unlabeled target nucleic acids. The method involves C sample having unlabeled target nucleic acids. The method involves attached to it in an array for the detection of multiple different target nucleic acids, the detection of multiple different target nucleic acids, to both providing nanoparticles having oligonucleotides bound to it, the Oligonucleotides bound to the annoparticles having a sequence that is complementary to at least a portion of the oligonucleotide bound to contacting the sample, the substrate, and the nanoparticles, the contacting occurring under conditions for hybridization of the target nucleic acids to the contacting a detectable change. The target cDNAs, c nanoparticles and substrate are contacted simultaneously under conditions effective for hybridization of the target cDNAs with the oligonucleotides bound to the substrate are contacted simultaneously under conditions effective for hybridization of the target cDNAs with the oligonucleotides bound to the substrate. The nanoparticles and with the capture nucleic acid sequences comprise an oligonucleotide, cDNA, or genomic condition in a sample of unlabeled target nucleic acid sequences comprise an oligonucleotide, cDNA, or genomic condition. The present sequence represents a detection probe that can condition and incompanies. The problems associated with fluorescent labeling and target can amplification. The present sequences are presented and target can advanced to detect target RNA sequences. The probe companies as a terroid and advanced and advanced to detect target RNA sequences. The probe companies as a terroid and advanced conditions and target can be used to detect target RNA sequences. The probe companies as a terroid and advanced conditions as a person of the condition and target can be used to detect target RNA sequences. The probe companies as a person of the condition and advanced condition and advanced conditions. The proper companies and advanc
                                                                                                                                                                                                                             nucleic acids by contacting the sample, a substrate having capture nucleic acid sequences and nanoparticles having bound oligonucleotides
                                                                                                                                                                                                        Detecting/quantifying gene expression in a sample of unlabeled target
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disulfide linker at the 5'-end followed by a recognition sequence.
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                       The invention relates to an oligonucleotide molecule (ON) labeled with several fluorophores of one or more types embedded in its backbone, where one or more of the fluorophores is not located at either the 3' or 5' terminus of ON. ON is useful for sequencing nucleic molecules. ON is useful for detecting biological molecules e.g., antibody, antigen, avidin, protein, peptide, bacteria, virus, blood cell or hormone. ON is capable of providing strong fluorescence signals at different wavelengths. This sequence corresponds to an example of an oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         el oligonucleotide molecule labeled with several fluorophores, useful detecting biological molecules e.g., antibody, antigen, avidin or
                                                                                                       Novel oligonucleotide molecule labeled with several fluorophores, useful for detecting biological molecules e.g., antibody, antigen, avidin or
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.00.0%; Pred. No. 6.8e+02;
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   (MONS ) MONSANTO TECHNOLOGY LLC.
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Matches 20; Conservative
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                                                                        WPI; 2005-063191/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Discriminating responsiveness of individual to taxanes, by detecting expression of 10 or more genes e.g., L43578, M33882, U32944 in sample derived from individual and determining responsiveness of sample based on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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capable of providing strong fluorescence signals at different wavelengths. This sequence corresponds to an example of an oligonucleotide of the invention.
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100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
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                                                                                                            Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                              O.7%; Score 20; DB 1; Lv. Local Similarity 100.0%; Pred. No. 6.8e+02; nes 20; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-2004; 2004WO-JP009692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2003; 2003JP-00270176.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Koizumi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TAIS ) TAISHO PHARM CO LTD (KATO/) KATO K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADW44957 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-APR-2005 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2005-101490/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NOGU/) NOGUCHI S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection results.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2005003352-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                  626
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The present invention relates to a method (M1) for stimulating proliferation of melanocyte cells in a human. (M1) involves administering or stem Cell Factor (SCF) protefail, or its biologically active fragments that stimulates growth of melanocyte cells, and optionally a carrier, to the human. The SCF is covalently conjugated to a water soluble polymer of the human. The SCF is covalently conjugated to a water soluble polymer of the proderings solved as early hematopoietic progenitor cells that primitive progenitors such as early hematopoietic progenitor cells that primitive progenitors such as early hematopoietic stem cells that are capable of maturing to erythroid, megakaryocyte, granulocyte, lymphocyte and macrophage cells, and non-hematopoietic stem cells such as neural stem cells and primordial garm stem cells. (M1) is useful in accelerating bone marrow regeneration, and in augmenting T cell contacted by a reduction in functional marrow mass due to toxic, radiant or immunological injury. (M1) is useful in treating ADDS, aplastic anemia, parroyemal nocturnal hemoglobinaria, myelogibioposis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     myelosclerosis, osteoperrosis, metastatic carcinoma, acute lenkemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, Niemann Pick disease, congestive splenomegaly, Kalaazar, sarcoidosis, primary splenic pancytopenia, disseminated fungus disease, fulminating septicemia, malaria, vitamin B12 and folic acid deficiency disease, pridoxine deficiency disease, and hypoigmentation discorders such as piebaldism and vitiligo. (M1) is useful in treating infertility states, intestinal damage resulting from irradiation or chemotherapy, and stem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stimulating proliferation of melanocyte cells in human, involves administering stem cell factor polypeptide or its biologically active fragments stimulating growth of melanocyte cells, and optionally carrier,
                                                                                                                                                                                                  CNS-Gen.; Antiparasitic; Antibacterial; Immunosuppressive;
Antinflammatory; Fungicide; Antifertility; AIDS; aplastic anemia;
paroxysemal nocturnal hemoglobinuria; osteopetrosis; acute leukemia;
multiple myeloma; hodgkins disease; lymphoma; gauchers disease;
niemann pick disease; sarcoidosis; plasmodium infection;
vitamin deficiency; hypopigmentation; vitiligo; infertility;
chronic myelocytic leukemia; cell proliferation; Stem Cell Factor; PCR;
                                                                                                                                                                                 Antiemetic; Cytostatic; Anti-HIV; Cardiovascular-Gen.;
                                                                                                                                       Universal Stem Cell Factor PCR primer 220-7, SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 33; 212pp; English.
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                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
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                   ADW93078 standard; DNA; 20
                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-160562/17.
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10-APR-1991;
25-NOV-1992;
                                                                                                 21-APR-2005
                                                                                                                                                                               Antianemic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-FEB-2005.
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24-AUG-1990
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                                                                                                                                                                                                                                                                                                                                                primer; ss
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                                                           ADW93078;
ADW93078/C
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                 primary thrombocythemia and acute leukemia. (M1) is useful in expanding searly hematopoietic progenitors in syngencic, allogencic or autoblogous bone marrow transplantation, and in enhancing the efficacy of genetherapy. The present sequence is a PCR primer used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid, e.g. 1-Aminocyclopropane-1-Carboxylate oxidase or ethylene insensitive, useful for producing green leaves in maize plants.
cell myeloproliferative disorders such as chronic myelogenous leukemia,
                                                                                                                                                                                     Gaps
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                                                                                                                                               0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
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100.0%; Pred. No. 6.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                Seguence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genetic engineering; transgenic plant; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   dT(20) primer used in cDNA synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; SEQ ID NO 49; 63pp; English
                                                                                                                                                                                                                2707 CTAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                   20 CTAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JUN-2004; 2004US-00876086.
                                                                                                                                                                                                                                                                                                                                ADY86103 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADZ47530 standard; DNA; 20
                                                                                 invention for cloning SCF.
                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                               20; Conservative
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Les 20; Conservative
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                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-241338/25.
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                                                                                                                                                                                                                                                                                                                                                                                                  02-JUN-2005
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                                                                                                                                                                                                                                                                                                                                                                   ADY86103;
                                                                                                                                               Query Match
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                                                                                                                                                                                 Matches
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Matches
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88888888
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Query Match 0.7%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 6.8e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                 89US-00422383.
90US-00537198.
90US-0053616.
90US-00589701.
91US-00684535.
93US-00982255.
93US-00172329.
93US-0017329.
                                                                                                                                      16-JUL-2003; 2003US-00620642
                                                                                                                                                                                                               19-JUN-2002; 2002US-00175608
                   30-JUN-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                              rat and human SCF cDNA.
                                                                                                                                                                                                                                                               Zsebo KM, Bosselman RA,
                                                                                                                                                                                                                            (ZSEB/) ZSEBO K M.
(BOSS/) BOSSELMAN R A.
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
                                                                                                                                                                                                                                                                            WPI; 2005-321855/33.
                                                                                                           US2005080250-A1
                                                                                                                                                                       01-OCT-1990;
10-APR-1991;
25-NOV-1992;
                                                                                             Unidentified
                                                                                                                                                                                           21-DEC-1993;
07-JUN-1995;
                                                                                                                                                                                                         07-AUG-2000;
                                                                                                                                                                24-AUG-1990;
                                                                                                                        14-APR-2005
                                                                                 primer; 88.
     ADZ47530
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The present invention relates to a method of stimulating growth of stromal cells in a human. The method involves administering to the human are cell factor (SCP) polypeptide and coptionally a pharmaceutically acceptable carrier. The invention is useful for treating immune disorders such as (acquired immune deficiency syndrome, severe combined immunodeficiency), genetic disorder (such as niemann pick disease), hematological disease (such as multiple myeloma, nodgkins disease, spleen disease, anemia), cancers (such as multiple myeloma, lymphoma), neurological disease (such as niemann pick disease), infection (such as Leishmania donovani infection), hypopigmentation disorders (such as piebaldism and vittligo), in the treatment of infertility states and for treating intestinal damage resulting from irradiation or chemotherapy and inflammation (such as sarcoidosis). The invention is also useful in gene therapy and SCF therapy. The present construction is also useful in gene therapy and SCF therapy. The cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stimulating growth of stromal cells for treating AIDS or severe combined immunodeficiency states in human, primary splenic pancytopenia, miliary tuberculosis, by administering human stem cell factor polypeptide and carrier to human.
                                                                                                      Stem cell factor; cell growth; immune disorder; immunomodulator; genetic disorder; hematological disease; antianemic; cancer; cytostatic; neoplasm; neurological disease; neuroprotective; infection; antimicrobial; hypopigmentation; dermatological; infertility; antiinfertility; inflammation; antiinflammatory; gene therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suggs SV, Martin FH;
Universal PCR primer, 220-7, SEQ ID NO: 33.
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Amplifying unknown nucleotide sequence adjacent to known nucleotide sequence, involves performing primary amplification of unknown nucleotide sequence using DNA walking annealing control primer and first target-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a method of amplifying an unknown nucleotide sequence adjacent to a known nucleotide sequence, comprising performing a primary amplification of the unknown nucleotide sequence using a DNA walking annealing control primer and a first target-specific primer by performing a first- and second-stage amplification of the unknown nucleotide sequence. The invention also relates to a DNA walking annealing control primer for amplifying an unknown nucleotide sequence adjacent to a known sequence of a DNA walking annealing control primer. This sequence represents a synthetic RT-PCR primer of the invention.
                                                                                                                                                                                                                             DNA amplification; RT-PCR; primer; 88; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'Match 0.7%; Score 20; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 6.8e+02; es 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 27; 66pp; English.
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CTAAAAAAAAAAAAAAAA 1
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                                                                                      Вb
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                                                                                    AEA01023 standard; DNA; 20
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                                                                                                                                                          (first entry)
                                                                                                                                                                                             Synthetic RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               (SEEG-) SEEGENE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          specific primer.
                                                                                                                                                                                                                                                                                                     WO2005045073-A1
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                                                                                                                                                                                                                                                                   Synthetic.
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                                                                                                                         AEA01023;
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20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chun J;
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                                                   RESULT 63
AEA01023/
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Gaps ö ADZ98001;

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The invention relates to a method for modulating, in a host cell, a protein-protein interaction between a first protein which is PRAK (P38-cegulated/activated protein kinase or MAPKAPK5) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host call a protein-protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, juvenile chronic arthritis, myositis, Crohn's disease, gastritis, colitis, unflammatory disease, proteitis, pelvic inflammatory disease, systemic lupus erythematosus, rhinitis, colitis, inflammatory disease, systemic lupus erythematosus, rhinitis, conjunctivitis, solaritis, chronic inflammatory polyneuropathy, Tertiary Lyme disease, psoriais, dermatitis or eczema. In the exemplification of the present invention examples of antisense oligonucleotides specific to nucleic acids encoding individual proteins in tables I to 82 are provided in SEQ ID NOS:11-223 (ADZ97857-ADZ98069).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPK5) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bush A;
                 protein interaction; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cimbora D, Heichman K, Bartel P, Mauck K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 153; 296pp; English.
                                                                                                                                                                                                                                                                                                                        2001US-0276179P.
2001US-0277013P.
2001US-0307233P.
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2000US-0185056P.
2000US-00727384.
2000US-0255063P.
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2001US-0259572P.
                                                                                                                                                      20-OCT-2003; 2003US-00690276
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2002US-00035344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-MAR-2002; 2002US-00100503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2005-371623/38.
                                                                                   US2005112118-A1.
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19-MAR-2001;
23-JUL-2001;
14-DEC-2001;
21-DEC-2001;
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21-DEC-2000;
04-JAN-2001;
04-JAN-2001;
                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                               04-JAN-2002;
                                                                                                                    26-MAY-2005
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Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
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0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels Query Match 0.77 Best Local Similarity 100. Matches 20, Conservative

647 GTAGCCAACATGTCAGGGTG 666 gracccaacarcresecere 1 20 ઠે g

ADZ98001 standard; DNA; 20 BP. RESULT 632 ADZ98001/c ü

protein interaction; antisense oligonucleotide; ss. Human antisense oligonucleotide SEQ ID NO:155 99US-0168379P. 2000US-0185056P. 2000US-00727384. 21-DEC-2000; 2000US-0256986P. 04-JAN-2001; 2001US-0259571P. 04-JAN-2001; 2001US-0259572P. 15-MAR-2001, 2001US-0276179P. 19-MAR-2001, 2001US-0277013P. 23-JUL-2001, 2001US-0307233P. 14-DEC-2001, 2001US-00014814. 20-OCT-2003; 2003US-00690276 2000US-0255063P 21-DEC-2001; 2001US-00024599 04-JAN-2002; 2002US-00035343 04-JAN-2002; 2002US-00035344 2002US-00099924 18-MAR-2002; 2002US-00100503 (first entry) US2005112118-A1. 28-JUL-2005 14-DEC-2000; 01-DEC-2000; 26-MAY-2005 

Bush A; Mauck K, Bartel P, Cimbora D, Heichman K,

(MYRI-) MYRIAD GENETICS INC

WPI; 2005-371623/38.

Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPKS) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.

Disclosure; SEQ ID NO 155; 296pp; English.

The invention relates to a method for modulating, in a host cell, a protein interaction between a first protein which is PRAK (P38-c regulated/activated protein kinase or MAPRARS) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host cell a protein-protein interaction between a first protein which is PRAK and a second protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., aschma, rheumatorid arthritis, juvenile chronic arthritis, myositis, cloim's disease, gastritis, colitis, ulcerative colitis, inflammatory bowel disease, proteitis, pelvic conjunctivitis, scleritis, chronic inflammatory polyneuropathy, Tertiary lyme disease, psoriasis, dermatitis or eczema. In the exemplification of the present invention examples of antisense oligonucleotides specific to nucleic acids encoding individual proteins in tables 1 to 82 are provided in SEQ ID NOS:11-223 (ADZ97857-ADZ98069).

Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;

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Gaps ö

Gaps ö Query Match
0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels

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643 AGCAGTAGCCAACATGTCAG 662

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The invention relates to a method for modulating, in a host cell, a protein protein interaction between a first protein which is PRAK (P38-requlated/activated protein kinase or MAPRAPK5) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host call a protein-protein interaction between a first protein which is PRAK and a second protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., ashma, rheumatoid arthritis, juvenile chronic arthritis, myositis, Crohn's disease, gastritis, colitis, ulcerative colitis, inflammatory bowel disease, proctitis, pelvic inflammatory disease, systemic lugus erythematosus, rhinitis, conjunctivitis, scleritis, chronic inflammatory polyneuropathy, Tertiary Lyme disease, psoriasis, dermatitis or eczema. In the exemplification of the present invention examples of antisense oligonucleotides specific to nucleic acids encoding individual proteins in tables I to 82 are provided in SEQ ID NOS:11-223 (ADZ97857-ADZ98069).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPKS) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bush A;
                                                                                                                                                                                                                                                                                                                                                                                            protein interaction; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                             Human antisense oligonucleotide SEQ ID NO:154.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mauck K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 154; 296pp; English
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20 AGCAGTAGCCAACATGTCAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000US-0256966P.
2001US-0259571P.
2001US-0259572P.
2001US-0276179P.
2001US-0277013P.
2001US-0017913P.
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2000US-00727384.
2000US-0255063P.
                                                                                                                                                   ADZ98000 standard; DNA; 20 BP.
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2002US-00035343.
2002US-00035344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-00100503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-OCT-2003; 2003US-00690276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                     28-JUL-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Heichman K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002US-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2005-371623/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS2005112118-A1
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04-JAN-2001;
04-JAN-2001;
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23-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-MAY-2005
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                                                                                                                                                                                                               ADZ98000;
                                                                                           RESULT 633
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The invention relates to a novel method for analyzing a nucleic acid. The method involves performing amplification and individual detection of a nucleic acid in a chip having several reaction layers for accommodating samples and reagents, to enable specific hybridization of the target nucleic acid and portions capable of controlling temperature conditions for amplification and detection of the nucleic acid. The invention further comprises a nucleic acid analyzer consisting of the chip as mentioned in the method, and a temperature control unit for performing hybridization for performing nucleic acid detection. The method enables convenient, cost-effective and highly accurate analysis of a nucleic acid acid. This sequence represents an oligo used in the nucleic acid analyzing method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Analyzing nucleic acid by performing amplification and individual detection of nucleic acid in chip having reaction layers for accommodating sample and reagent, and portions capable of controlling temperature conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                    Gaps
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0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                            DNA detection; biochip; hybridization; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 1; 50pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HITA-) HITACHI HIGH TECHNOLOGIES CORP.
                                                                                                                                                                                                                                                                                         Nucleic analysis 20-mer Thymine oligo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2709 AAAAAAAAAAAAAAAAA 2728
                                                                    645 CAGTAGCCAACATGTCAGGG 664
                                                                                           CAGTAGCCAACATGTCAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 AAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8251/c
AEB28251 standard; DNA; 20 BP.
                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-DEC-2003; 2003WO-JP015490.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-DEC-2003; 2003WO-JP015490
                                                                                                                                                                                      AEA98126 standard; DNA; 20
                                                                                                                                                                                                                                                          25-AUG-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kajiyama T, Takahashi S;
                                    20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2005-425411/43.
                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                               WO2005054458-A1
                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUN-2005.
                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                          AEA98126;
                                                                                                       20
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       Query Match
Best Local (
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                                      Matches
                                                                                                                                                          RESULT 634
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ID AEB28
XX
                                                                                                                                                                             AEA98126,
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Sequence 20 BP; 3 A; 6 C; 5 G; 6 T; 0 U; 0 Other;

AEB28251;

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This invention represents a novel cosmetic or pharmaceutical composition for treating epithelial covering tissue which comprises superstructure-forming nucleic acid sequences. The composition can also be used in fabric softeners, hand-washing products. Dody and hair care products, hair dyes or manual dishwashing products. The superstructures are G quadruplexes frayed wires or i motife. The sequences are 30.40 nucleotides long, have five or more C, G or I nucleotides in tandem, no CpG motifs, no nonmethylated CG dinucleotides, are poly1, polyC or polyG homopolymers and are optionally modified by replacing phosphodiester inkages with methylphosphonate, phosphoramidate, phosphorothicate or pertopyranoses or 3, 5, -canbocyclically bridged 2'-deoxyribose pertopyranoses or 3, 5, -canbocyclically bridged 2'-deoxyribose composition of the invention inhibits the release of IL-8. The composition is useful for preventing or treating inflammatory changes to epithelial covering tissue, including changes caused by pathogens, cutoimmune reactions, tumor necrosis factor, toxins and irritants, especially aging processes, psoriasis, atopic eczema, dry skin, alopecia areata, vitiligo, bullous diseases, rajection reactions, sunburn and parodontosis. This sequence represents a phosphorothicate or phosphodiester oligonucleotide used to illustrate the method of the
                                                                                                                                         dermatological disease; antiinflammatory; antiallergic; aging; eczema; alopecia; epidermolygis bullosa; graft rejection; periodontal disease; psoriasis; antipsoriatic; unburn; vitiligo; inflammation; detergent; dye; pigment; ss; primer; phosphorothioate; phosphodiester.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cosmetic or pharmaceutical composition for treating epithelial tissue comprises superstructure-forming nucleic acid sequences.
                                                                                                                      pharmaceutical; skin allergy; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 2; 71pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-DEC-2004; 2004WO-EP014195.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHEN-) PHENION GMBH & CO KG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-2003; 2003DE-01061502
                                       22-SEP-2005 (first entry)
                                                                               Oligonucleotide 100T-PTO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-512612/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kippenberger S,
                                                                                                                                                                                                                                                                                                                                                                                                          WO2005063300-A2
                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                         cosmetics;
                                                                                                                                                                                                                                           Synthetic
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Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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2709 AAAAAAAAAAAAAAAA 2728 100.08; FIL AAAAAAAAAAAAAAA 1 20

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dermatological disease; antiinflammatory; antiallergic; aging; eczema; alopecia; epidermolysis bullosa; graft rejection; periodontal disease; psoriasis; antipsoriatic; unburn; vitiligo; inflammation; detergent; dye; pigment; 88; primer; phosphorothioate; phosphodiester.
                                                                                                                                                            /note= "phosphorothioate or phosphodiester linkages"
                                                                      cosmetics; pharmaceutical; skin allergy; dermatological;
                                                                                                                              Location/Qualifiers
                                                                                                                                                    /mod_base= OTHER
                                                                                                                                                                                                     14-DEC-2004; 2004WO-EP014195.
             AEB28250 standard; DNA; 20
                                         (first entry)
                                                                                                                                              ø
                                                        Oligonucleotide 100A-PTO.
                                                                                                                                              tag=
                                                                                                                                                                         WO2005063300-A2
                                                                                                                             Key
modified_base
                                          22-SEP-2005
                                                                                                                                                                                       14-JUL-2005.
                                                                                                                Synthetic
                           AEB28250;
RESULT 636
        AEB28250
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/mod\_base= OTHER /note= "phosphorothioate or phosphodiester linkages"

Location/Qualifiers

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\*tag=

Bernd A, Kippenberger S, Kaufmann R, (PHEN-) PHENION GMBH & CO KG.

epithelial covering

Bock A;

Bernd A,

Kaufmann R,

23-DEC-2003; 2003DE-01061502.

Bock A;

WPI; 2005-512612/52.

Cosmetic or pharmaceutical composition for treating epithelial covering tissue comprises superstructure-forming nucleic acid sequences.

Disclosure; SEQ ID NO 1; 71pp; German.

This invention represents a novel cosmetic or pharmaceutical composition for treating epithelial covering tissue which comprises superstructureforming nucleic acid sequences. The composition can also be used in fabric softeners, hand-washing products, body and hair care products, hair dyes or manual dishwashing products. The superstructures are G quadruplexes, frayed wires or i motifs. The superstructures are G quadruplexes, frayed wire or mere C, G or I nucleocides in tandem, no composition, have five or mere C, G or I nucleocides in tandem, no CG motifs, no nonmethylated CG dinucleotides, are poly!, polyC or polyG nucleotides and are optionally modified by replacing phosphodisater inhages with methylphosphonate, phosphoramidate, phosphorothicate or hydroxylamine linkages, by replaced ribose with other hexo- or pentopyranoses or 3, 5, -carbocyclically bridged 2'-deoxyribose cervices or an expensive composition of the invention inhibits the release of IL-8. The composition is useful for preventing or treating inflammatory changes to epithelial covering tissue, including changes caused by pathogens, cuttoimmune reactions, tumor necrosis factor, toxins and irritants, especially aging processes, peoriasis, atopic eczema, dry skin, alogecia especially aging processes, psoriasis, atopic eczema, dry skin, alope areata, vitiligo, bullous diseases, rejection reactions, sunburn and parodontosis. This sequence represents a phosphorothioate or phosphodiester oligomucleotide used to illustrate the method of the invention.

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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Gaps

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0; Indels

Length 20;

0.7%; Score 20; DB 1; Le 100.0%; Pred. No. 6.8e+02; lve 0; Mismatches 0;

Conservative

Local Similarity les 20; Conserv

Query Match Best Loca Matches

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Gaps
                                           ö
0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels
 Query Match
Best Local Similarity 100.
Matches 20; Conservative
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2709 AAAAAAAAAAAAAAAA 2728

8

1 AAAAAAAAAAAAAAAA 20

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RESULT 638

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The invention relates to a novel method for preparing an oligonucleotide with phosphorothioate internucleoside linkage(s), by phosphitylating the 5.-hydroxyl of a nucleic acid group to form a phosphite intermediate; and oxidizing the phosphite intermediate with a substituted acetyl disulfide or acetyl disulfide in an aprotic solvent with a protic solvent or a basic solvent enough to convert the phosphite intermediate to a phosphorothioate. The method is useful in molecular biological research and in applications such as anti-viral therapy. Modified oligonucleotides are also useful as antisense reagents because they can block ribonucleic acid translation, and are nuclease resistant. Phosphorothioate-containing oligonucleotides are also useful in determining the stereochemical pathways of certain enzymes that recognize nucleic acids. The sequences shown in (ABC31667-ABC31682) represent oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Preparation of oligonucleotide with phosphorothioate internucleoside linkage(s) that determines stereochemical pathways of nucleic acid recognizing enzyme by, phosphitylating 5'-hydroxyl of nucleic acid and oxidizing phosphite intermediate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                             note= "OTHER=phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "OTHER=2'-O-methoxyethyl-thymine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                    Phosphorothioate oligonucleotide SEQ ID NO:10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cheruvallath ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 12; SEQ ID NO 10; 12pp; English.
                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                             /mod_base= OTHER
                                                                                                                                                                       phosphorothioate; antiviral; ss.
                              BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAY-2004; 2004US-00847502.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-JAN-2001; 2001WO-US000715.
12-DEC-2002; 2002US-00181200.
                              AEC31676 standard; DNA; 20
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/*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.77
Best Local Similarity 100.
Matches 20, Conservative
                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                 *tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cole DL, Ravikumar VT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2005-615154/63
                                                                                                                                                                                                                                                                                                                                                                                                                      US2005165226-A1
                                                                                                                                                                                                                                              Key
modified_base
                                                                                                                                                                                                                                                                                                                                 modified base
                                                                                                   20-OCT-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JUL-2005
                                                                                                                                                                                                             Synthetic.
                                                                 AEC31676;
RESULT 637
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The invention relates to a novel method for preparing an oligonucleotide with phosphorothioate internucleoside linkage(s), by phosphitylating the 5'-hydroxyl of a nucleic acid group to form a phosphite intermediate; and oxidizing the phosphite intermediate with a substituted acetyl disulfide or acetyl disulfide in an aprofic solvent with a protic solvent or a basic solvent enough to convert the phosphite intermediate to phosphorothioate. The method is useful in molecular biological research and in applications such as anti-viral therapy. Modified oligonucleotides are also useful as antisense reagents because they can block ribonucleic acid translation, and are nuclease resistant. Phosphorothioate-containing oligonucleotides are also useful in determining the stereochemical pathways of certain enzymes that recognize nucleic acids. The sequences shown in (AEC31667-AEC31682) represent oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preparation of oligonuclectide with phosphorothicate internucleoside linkage(s) that determines stereochemical pathways of nucleic acid recognizing enzyme by, phosphitylating 5'-hydroxyl of nucleic acid oxidizing phosphite intermediate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gapa
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                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "OTHER=phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                       /note= "OTHER=2'-O-methylthymine"
                                                                                                         Phosphorothicate oligonuclectide SEQ ID NO:15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cheruvallath ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 22; SEQ ID NO 15; 12pp; English.
                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
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                                                                                                                                         phosphorothioate; antiviral; ss.
           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JAN-2001; 2001WO-US000715.
12-DEC-2002; 2002US-00181200.
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Best Local Similarity 100.0
Matches 20, Conservative
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                                                                          (first entry)
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             AEC31681 standard; DNA;
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-615154/63.
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modified_base
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                                                                            20-0CT-2005
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                                                                                                                                                                            Synthetic.
                                            AEC31681;
AEC31681,
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Gaps

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2709 AAAAAAAAAAAAAAAA 2728

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20 AAAAAAAAAAAAAAA 1

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microarray, and assaying the sample for antimicrobial activity. Host cells comprising the nucleic acid may be used to obtain the claimed polypeptide. An antibody which binds the claimed polypeptide an also be used to identify an antimicrobial protein. A claimed pharmaceutical composition comprising the antimicrobial polypeptide and/or nucleic acid
                                                                                                                                                                                                                                  Example 1; SEQ ID NO 10; 84pp; English
            AEC37012 standard; DNA; 20
                                       (first entry)
                                                   Oligodeoxythymidine dT20.
                                                                                                                                                                                           WPI; 2005-582941/59
                                                                             drug screening; ss
                                                                                                       WO2005079523-A2.
                                      03-NOV-2005
                                                                                                                    01-SEP-2005
                                                                                                                                                                              Evans DL,
                                                                                          Synthetic.
                          AEC37012;
RESULT 639
      AEC37012/
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RESULT 640 AEC37011 888888888 ઠ g Antimicrobial; antibacterial; fungicide; protozoacide; bacterial infection; gene therapy; non-toll like New teleost-derived antimicrobial non-scavenger Receptor A, non-toll l receptor polypeptide, useful for treating a disorder resulting from a microbial infection and/or reducing antibiotic resistance. Praveen K; Leary JH, Kaur H, Jaso-Friedmann L, (UYGE-) UNIV GEORGIA RES FOUND INC. BP. 18-FEB-2005; 2005WO-US005398. 18-FEB-2004; 2004US-0545370P 01-NOV-2004; 2004US-0623909P

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Gaps

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0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; ive 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAAAA 2728

Matches 20; Conservative

Similarity

Query Match Best Local

20 AAAAAAAAAAAAAAA 1

AEC37011 standard; DNA; 20 BP.

(first entry)

03-NOV-2005

AEC37011;

Oligodeoxyadenosine dG20.

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

reduce antibiotic resistance. The present sequence is that of oligodeoxythymidine dT20. In an example from the invention, dT20 was used to identify membrane proteins on teleost nonspecific cytotoxic cells that bind single base oligodeoxynucleotide ligands.

to reduce antibiotic resistance. The polypeptide is present in an amount effective to inhibit microbial growth, e.g. bacterial, protozoa or fungal growth in a subject, e.g. a mammal (human), or in an amount sufficient to

is used to treat a disorder resulting from a microbial infection and/or to reduce antibiotic resistance. The polypeptide is present in an amount

Antimicrobial; antibacterial; fungicide; protozoacide; bacterial infection; fungal infection; protozoal infection; gene therapy; /note= "OTHER= optional 5' biotin label" Location/Qualifiers (UYGE-) UNIV GEORGIA RES FOUND INC. /mod base= OTHER 18-FEB-2004; 2004US-0545370P. 01-NOV-2004; 2004US-0623909P. 18-FEB-2005; 2005WO-US005398. \*tag= a drug screening; ss. WPI; 2005-582941/59 WO2005079523-A2 Key modified\_base Evans DL, Synthetic An non-toll like receptor polypeptide having a molecular weight of about 22-30 kba and having properties selected from: (a) (i) being obtainable from a telecost, e.g. catfish (Ictalurus punctatus), mammalian monocytess or mammalian manocytess or mammalian monocytess or mammalian monocytes or monocytes or monocytes or monocytes or monocytes or monocytes or mammalian monoc The invention provides an isolated antimicrobial non-scavenger receptor

New teleost-derived antimicrobial non-scavenger Receptor A, non-toll like receptor polypeptide, useful for treating a disorder resulting from a microbial infection and/or reducing antibiotic resistance. Praveen K; Leary JH, Kaur H, Jaso-Friedmann L,

Example 1; SEQ ID NO 9; 84pp; English.

A, non-toll like receptor polypeptide having a molecular weight of about 22-30 kDa and having properties selected from: (a) (i) being obtainable from a teleost, e.g. caffish (Ircalurus punctatus), mammalian monocytes or mammalian macrophages, (ii) binds to oligoguanosine, (iii) comprises 58 basic amino acids selected Lys and Arg, (iv) comprises 50 hydrophobic comprises 50 polar amino acids selected from Ala, Ile, Leu, Phe, Trp and Val, and (v) comprises 50 polar amino acids selected from Amino Cys, Gln, Ser, Thr and Tyr, containing 11 Lys-rich motifs, (b) comprises an amino acid sequence selected from: amino acid residues 1-60, 1-118, 27-51, 136-159, or 173-The invention provides an isolated antimicrobial non-scavenger receptor

CC 203 of the catfish nonspecific cytotoxic cells antimicrobial protein-1 (CC CATCHS NCAMP-1; (d) an allelic variant of catfish NCAMP-1; (e) a polypeptide encoded by a nucleic acid molecule that hybridizes under stringent conditions to the opposite strand of a catfish NCAMP-1; (e) a polypeptide encoded by a nucleic acid molecule catfish NCAMP-1; (n) of a catfish NCAMP-1 uncleic acid molecule AEC37006; (f) catfish NCAMP-1 comprising one acid substitutions; and (g) a fragment of catfitsh NCAMP-1 uncleic acid molecule AEC37006; (f) catfish NCAMP-1 uncleic acid molecule AEC37006; (f) catfish NCAMP-1 comprising one or more of these polypeptides is activity. A library compounds with the polypeptide comprises contacting candidate compounds with the polypeptide or library and selecting those capable of inhibiting the bioactivity of the polypeptide contacting consisting of NCCs from a teleost fish; mammalian monocytes or mammalian macrophages; isolating membranes from cultured cells consisting of NCCs from a teleost fish; membranes from cultured cells consisting of NCCs from a teleost fish; prolypeptide binds to oligoguanosine and/or has antimicrobial activity. Also claimed are nucleic acids encoding the antimicrobial polypeptide binds to oligoguanosine and or membranes; and determining if the colypeptide binds to claimed physicial public acids. A claimed method for detecting the presence or absence or absence of an oligoguanosine of another acids. A claimed method for detecting the presence or absence or absence or absence or along the antimicrobial activity. Host carcoarray, and assaying the sample for antimicrobial activity. Host call mind method for actecting the presence or absence or absence or colypeptide. An antibody which binds the claimed polypeptide can also be used to treat a disorder resulting from a microbial infection and/or composition composition composition composition protein and an amount sufficient to reduce antibiotic resistance. The present sequence is that of criedic and another and an electric a 

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The invention relates to a novel reaction detection chip e.g. a DNA chip for diagnosis of disease. The novel chip has porous-glass particles that are embedded in organic film (e.g. vinyl acetate) as a monolayer on a substrate (e.g. silicon). The analyte is preferably detected by fluorescence. The invention further includes a process for the manufacturing of the detection chip. The detection chip is useful e.g. as a DNA chip for diagnosis of cancer, and albo for analysis of medical treatment of pathogenesis related gene of multifactorial disorder. This oligo sequence represents a control DNA probe used in the novel detection chip of the invention.

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

Reaction detection chip e.g. deoxyribonucleic acid chip for diagnosis of cancer, has porous-glass particles that are embedded in organic film, as

SEQ ID NO 7; 19pp; Japanese.

monolayer.

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ;; DB 1; Length 20; 6.8e+02; Indels ö 0.7%; Score 20; DB 100.0%; Pred. No. 6.8 ative 0; Mismatches Query Match
Best Local Similarity 100.
Matches 20; Conservative

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AEC91079 standard; DNA; 20 AEC91079; RESULT 641 AEC91079/ 

17-NOV-2005 (first entry)

p53 cancer inhibition gene negative control DNA probe, SEQ ID 7. biochip; probe; ss.

Synthetic.

JP2005249429-A.

01-MAR-2004; 2004JP-00056758 15-SEP-2005

01-MAR-2004; 2004JP-00056758

(EBAR ) EBARA CORP.

The present invention relates to a method (M1) for labeling double stranded nucleic acid for efficient detection of DNA or RNA. The method comprises using a carbodismide derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid. (M1) is useful for labeling double stranded or single stranded nucleic acid or detecting single nucleotide polymorphisms. The present sequence was used to illustrate the method of the invention.

Example 1; Page 24; 40pp; Japanese.

nucleic acid

Ogure N;

Nakamura K, Abe Y, WPI; 2005-668391/69

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Labeling double stranded nucleic acid, involves utilizing carbodismide derivative for labeling thymine, uracil and guanine, which exists in mismatch region of nucleic acid or unstable region of hydrogen bond of
                                                                                                                                                                            Oligonucleotide ODN1 used to illustrate nucleic acid labeling method.
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  Length 20;
                    0; Indels
0.7%; Score 20; DB 1; L
100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0;
                                                                                                                                                                                                  DNA detection; RNA detection; SNP detection; 88.
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                                                                                                                  AED13293 standard; DNA; 20
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            Best Local Similarity 100.
Matches 20; Conservative
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    Query Match
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AED13293
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(first entry)

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                                                                                                                                                                                                                                                                                                                                                                        Molecule separation for biotechnology, comprises forming a nonturbulent flow condition for each solute molecule based on thermodynamic characteristics and changing flow conditions at arbitrary points.
                                           Gaps
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                          Length 20;
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                                           Indels
      Sequence 20 BP; 19 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
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ches 0;
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                                                                                                                                                                                                      molecule separation; purification; probe; ss.
                        0.7%; Score 20; DB
100.0%; Pred. No. 6.8
:ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                      Miyazaki M,
                                                            2708 TAAAAAAAAAAAAAAA 2727
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                                                                                                                          AED11364 standard; DNA; 20 BP
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                                 3est_Local Similarity 100.
fatches 20; Conservative
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                       Query Match
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AED42120 standard; DNA; 20 BP AED42120 RESULT 644 AED42120/ SXB

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Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive; Antidiabetic; Cardiant; Neuroprotective; Antiasthmatic; Antinflammatory; Antibacterial; Osteopathic; Anorectic; Virucide; Anti-HIV; Nephrotropic; Antiatteriosclerotic; Muscular-Gen.; protein interaction; protein microarray; cancer; familial adenomatous polyposis; protein microarray; cancer; familial adenomatous polyposis; gastrointestinal-gen.; ischemia; cerebrovascular ischemia; autoimmune disease; heart disease; neuroimmune disease; heart disease; neurodegenerative disease; asthma; inflammation; sepsis; osteoporosis; obesity; viral infection; acquired immune deficiency syndrome; glomerulonephritis; atherosclerosis; muscular dystrophy; antisense oligonucleotide; ss.
Antisense oligo of human protein-protein complex polypeptide, SEQ ID 186
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30-APR-2001; 2001US-0287513P.
10-UUL-2001; 2001US-0304101P.
23-UUL-2001; 2001US-030733P.
22-OCT-2001; 2001US-034381P.
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16-APR-2001; 2001US-0284095P.
17-APR-2001; 2001US-0284404P.
17-APR-2001; 2001US-0284404P.
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06-FEB-2002;
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WPI; 2005-664172/68.

second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma. New isolated protein complex having a first protein interacting with a

Disclosure; SEQ ID NO 186; 198pp; English.

The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method for selecting modulators of the protein complex; a method of selecting modulators of an interaction between a first protein and a second protein; and the treating and/or preventing of diseases and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders associated with the protein complexes. The diseases and disorders include

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                                                                                                                                                                                                                                                                                                                                                                                            Antisense oligo of human protein-protein complex polypeptide, SEQ ID 187.
cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, osteoporosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an antisense oligo of a human protein which forms part of a protein-protein complex of the invention.
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                                                                                                                                     DB 1; Length 20; 6.8e+02;
                                                                                                           Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                  100.0%; Prec. ...
                                                                                                                                        0.7%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                   648 TAGCCAACATGTCAGGGTGG 667
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2001US - 0284095P
2001US - 028420P
2001US - 0284204P
2001US - 0287513P
2001US - 0304101P
2001US - 0304101P
2001US - 0304389P
2001US - 0343818P
2001US - 0343818P
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2001US-0276259P.
2001US-0276179P.
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hes 20; Conservative
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22-OCT-2001;
25-OCT-2001;
04-JAN-2002;
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-JAN-2002;
-FEB-2002;
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17-APR-2001;
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30-APR-2001;
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The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method of compulses: a protein microarray complex; a method of selecting modulators of the protein complex; a method of selecting compulses of an interaction betweenting of diseases and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders complexes, and in treating and/or preventing diseases and disorders include cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, cinflammatory disorders, sepsis, osteoporosis, obesity, viral infection, albo, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an antisease oligo of a human protein which forms
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                                                                                                                                                                                                                                          New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia,
                                                                                                                                                                                                                                                                        second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma
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                                                                                              Wettstein DA, Heichman
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Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 187; 198pp; English.
                                                                                                      Sugiyama J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  665
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04-JAN-2001; 2001US-0259573P.
14-MAR-2001; 2001US-0276259P.
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                              (MYRI-) MYRIAD GENETICS INC.
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                                                                                                  Bartel P, Cimbora D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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Matches
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The present invention relates to a method for augmenting T-helper 1 cells
                     Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiuler; Dermatological; Antiallergic; helper T.lymphocyte; immune stimulation; inflammatory psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; mimune stimulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy; contact dermatitis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inhibit prostagiandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating non-allergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
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Pred. No. 6.8e+02;
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                                                                                                                                                                                                            /mod_base= OTHER
/note= "Phosphorothioate backbone"
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                                                                                       contact dermatitis; ss; phosphorothioate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               contact dermatitis or latex dermatitis.
                                                                                                                                                            Location/Qualifiers
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/*tag=
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Best Local Similarity
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                                                                                                                                                                          modified base
                                                                                                                                                                                                                                                                                               10-NOV-2005
                                                                                                                        Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method for selecting modulators of the protein complex; a method of selecting modulators of an interaction between a first protein and a second protein; and the treating and/or preventing of diseases and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders associated with the protein complexes. The diseases and disorders include cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsies, osteoporosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an antisense oligo of a human protein which forms
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 185; 198pp; English.
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                                                                            2001US-028324P
2001US-0287113P
2001US-0304101P
2001US-0304223P
2001US-03438BP
2001US-03438BP
2002US-034384P
2002US-034384P
2002US-034384P
2002US-034384P
                               2001US-0284095P.
2001US-0284220P.
2001US-0284404P.
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Best Local Similarity 100."
Matches 20, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               Bartel P, Cimbora D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2005-664172/68.
                                                                                                                               23-JUL-2001;
22-OCT-2001;
25-OCT-2001;
04-JAN-2002;
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06-FEB-2002;
14-MAR-2002;
14-MAR-2002;
18-MAR-2002;
                                                                                 19-APR-2001;
30-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                               29-APR-2002;
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                                                                                                                     10-JUL-2001;
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Gaps

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The present invention relates to a method for augmenting T-helper 1 cells administrating an immunostimulatory nucleic acid (1) to induce Thl-like immune activation in a subject. The method comprises immune activation, and administrating a cyclooxygenase inhibitor Illike inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (1) is useful for treating nonallergic inflammatory diseases such as psoriacis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides a method for detecting the presence or absence of a single target molecule or target analyte (e.g. nucleic acid, protein, lipid, bacterium). The method involves contacting sample with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light scattering complex formed.
                                                     Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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100.0%; Pred. No. 6.8e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                       Disclosure; SEQ ID NO 537; 58pp; English
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                                                                                                                                                                                                                                                                               contact dermatitis or latex dermatitis.
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04-NOV-2003; 2003US-0517450P.
03-MAY-2004; 2004US-0567874P.
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Best Local Similarity 100.
Warches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NANO-) NANOSPHERE INC
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                           WPI; 2005-768014/78.
Berg DJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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Krieg AM,
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                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (1) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one buch immunostimulatory nucleic acid. (I) is useful for treating non-allergic inflammatory diseases such as psoriasis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunostimulant, Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; immune stimulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy;
                                                                                                                                                                                                                                                                   Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                  to augment T-helper1 cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 533; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2709 AAAAAAAAAAAAAAAA 2728
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29-MAR-2002; 2002US-00112653.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               L2-MAY-2005; 2005US-00127654
                                                                                                                          29-MAR-2001; 2001US-0279642P
29-MAR-2002; 2002US-00112653
                                                                                             12-MAY-2005; 2005US-00127654
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      contact dermatitis; ss
                                                                                                                                                                                                                                    WPI; 2005-768014/78
                                                                                                                                                                                                       Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                US2005250726-A1
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                                                               10-NOV-2005
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Synthetic
                                                                                                                                                                                                       Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AED75401;
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one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is an oligonucleotide used to prepare aptamer-coated gold probe arrays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides a method for detecting the presence or absence of a single target molecule or target analyte (e.g. nucleic acid, protein, lipid, bacterium). The method involves contacting sample with one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is a T20 diluent which is used in the preparation of aptamer-coated gold probes. Note: The present sequence is the SBQ ID NO: 26 which is given in the sequence listing. This sequence differs from the property of the specification (see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light
                                                                                                                                                                                                                                                                                                                                                                                                                                  T20 diluent SEQ ID: 26 #1 used to prepare aptamer-coated gold probes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Garimella
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                                                                                                                                    0.7%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 6.8e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analyte detection; DNA detection; protein detection; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= Linked to a steroid"
                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 26; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                         20 AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                             AED67955 standard; DNA; 20 BP
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29-MUG-2003; 2003US-04990J4P.
04-NOV-2003; 2003US-0517450P.
03-NAY-2004; 2004US-0567874P.
27-MAY-2004; 2004US-0085484B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-2004; 2004US-00995051
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  scattering complex formed
                                                                                                                                                                        20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lucas A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2005-784662/80.
                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                   12-JAN-2006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                 AED67955;
                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                             RESULT 651
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light
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                                                                                                                                Gaps
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                                                            Score 20; DB 1; Length 20;
Pred. No. 6.8e+02;
                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 6.8e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Senical M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analyte detection; DNA detection; protein detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Deoxyadenosine spacer oligonucleotide SEQ ID NO: 31.
Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                             100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Muller UR,
                                                                                                                                                                                      2709 AAAAAAAAAAAAAAAAA 2728
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04-NOV-2003; 2003US-0517450P.
03-MAY-2004; 2004US-0567874P.
27-MAY-2004; 2004US-00854848.
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                                                                                                                                                                                                                                                                                                                                                                                                                  AED67960 standard; DNA; 20
                           Query Match
Best Local Similarity luv...
Best Local Similarity luv...
20; Conservative
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RESULT 653

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The invention relates to a purified antibody that is specifically immunoreactive with a stem cell factor (SCF) or SCF receptor. Also described: (1) a hybridoma cell line producing a monoclonal antibody that is specifically immunoreactive with a SCF protein; (2) inhibiting the activity of a mast cell population; (3) decreasing blood cell profleration or activity in a in a mammal; (4) decreasing the interaction between a SCF and an SCF receptor in a cell population; (5) treating a mammal having a disorder mediated through the interaction of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF with an SCF receptor; and (6) a pharmaceutical composition of polypeptide, and a pharmaceutically immunoreactive with an SCF receptor in a cell population, decreasing the interaction maturation or activity in a in a mammal, decreasing the interaction between a SCF and an SCF receptor in a cell population, and treating a mammal having a clivity of maturation or activity in a in a mammal, decreasing the interaction between a SCF and an SCF receptor in a cell population, and treating a mammal having a clivity of maturation or activity composition, and methods are useful for treating disorders and the antibody, composition, and methods are useful for treating a universal cuberculosib, or malaria. The present sequence represents a universal oligonaclecide sequence which can be used as a probe or a PCR primer in the amplification and sequence mand and human SCF, which is used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New stem cell factor antibody, useful for treating hematopoietic disorders such as anemia, leukemia, lymphoma, HIV, tuberculosis, or
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0.7%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.8e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                 antibody; stem cell factor; probe; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Martin FH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 33; 217pp; English.
                                                                                                                            Universal oligonucleotide SEQ ID NO:33.
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                                                                                                                                                                                                                                                                                                                                                     89US-00422383.
90US-00537198.
90US-00573616.
91US-00684535.
92US-00172329.
95US-00172329.
                 BP.
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                 AEE01383 standard; DNA; 20
                                                                                         (first entry)
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BOSSELMAN R A.
SUGGS S V.
MARTIN F H.
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10-APR-1991;
25-NOV-1992;
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                                                                                         26-JAN-2006
                                                                                                                                                                                                                                                                                 24-NOV-2005
                                                                                                                                                                                                                                                                                                                                                       16-OCT-1989;
11-JUN-1990;
                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zsebo KM,
                                                    AEE01383;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZSEB/)
(BOSS/)
(SUGG/)
(MART/)
AEE01383/c
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The present sequence is a negative control DNA probe used in the covalidation of the novel reactive detection biochip which is the subject of the invention. The present inventions relates to a novel reactive detection chip comprising spots formed by immobilizing several porous particles in a single-layers formed by immobilizing several porous particles in a single-layers formed by immobilizing several porous particles are transparent to incident light and have probe conscious bactuling molecules in a sample. In these chips, the number of probes and spot thickness can be stably controlled, and the three-confinements are stably increased, consequently improving the signal components in a spotter (which is claimed), hearting the substrate to substrate, applying the porous particles in a spotted arrangement on the confinement of immobilize the particles and removing excess of porous particles in the organic film (mapping the detection capacitics in chip manufacture confinement in maintains the dispersed state of porous particles without and any any additive to the spotting solution, and thus enabling the chip to be manufactured.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reactive detection chip e.g. DNA chip to recognize functional molecule in genetic diagnosis, comprises spots formed by immobilizing porous particles including probes, in single-layered state on substrate, with particles transparent to light.
                                                                                                                                                                                                                                                                                        ss; biochip; p53 gene; tumor suppressor gene p53; DNA detection; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "conjugated porous glass particle"
                                                                                                                                                                                                                                                      Poly-thymine negative control probe SEQ ID NO: 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 7; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
2707 CTAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/mod_base= OTHER
                                        20 CTAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAR-2004; 2004JP-00056758.
12-AUG-2004; 2004JP-00235221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ogure N
                                                                                                                                      AED85822 standard; DNA; 20
                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nakamura K, Abe M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2006-016978/02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (EBAR ) EBARA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                            Key
modified_base
                                                                                                                                                                                                                                                                                                                                    Synthetic.
Unidentified.
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                                                                                                    RESULT 654
                                                                                                                      AED85822/
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Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

manufactured more efficiently

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Gaps

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human or other subject. (M1) comprises: (a) obtaining hematopoiesis in a human or other subject. (M1) comprises: (a) obtaining hematopoietic brogenitor cells from the human or other subject; (b) expanding the cells obtained in step (a) by adding to the cell a stem cell factor (SCF) collypeptide having a 195, 208 or 245 amino acid sequence of AEB60706, AEB60708 or AEB60725, or its biological active fragments that stimulate growth of hematopoietic progenitor cells; and (c) administering to the human or other subject the expanded hematopoietic progenitor cells; obtained in step (b), therefore restoring hematopoiesis to effect hematopoiesis in the human or other subject. Also described is a method comprises; (M2) for expanding hematopoietic progenitor cells ex vivo, which comprises; (a) obtaining hematopoietic progenitor cells ex vivo, which comprises; (a) obtained in step (a) by adding to the cells the comprises; (a) obtained in step (b) aspanding the cells obtained in step (a) by adding to the cells the chancing hematopoietic progenitor cells, where the hematopoietic cells are chosen from stem cells, lymphoid progenitor cells, myeloif progenitor expanding hematopoietic progenitor cells, where the hematopoietic cells, are chosen from stem cells, lymphoid progenitor cells, myeloif progenitor cells, magakarycoytes and erythroblasts. (M1) is useful for treating various stem cell deficiencies such as aplastic anemia, parcoxymal nocturnal hemaglobinuria, myeloifbrosis, myelosclerosis, Gaucher's disease, Niemann-Pick disease, Hodgkin's disease, Kala-azar, sarcoidosis,
                                         ö
                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enhancing hematopoiesis in human, comprises expanding hematopoietic progenitor cells by adding stem cell factor polypeptide to cells and
                                         ö
     Length 20;
                                     0; Indels
                                                                                                                                                                                                                                                                                                          Universal stem cell factor PCR primer SEQ ID NO:33
DB 1; Ler
. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                hematopoiesis; stem cell factor; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Martin FH
 0.7%; Score 20; DB
100.0%; Pred. No. 6.8
/ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 33; 213pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            progenitor cells by adding stem cell fi
administering expanded cells to human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Suggs SV,
                                                                       2709 AAAAAAAAAAAAAAAA 2728
                                                                                             90US-00537198.
90US-00573616.
90US-00589701.
                                                                                                                                                                                                 AEE60695 standard; DNA; 20 BP
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92US-00982255.
93US-00172329.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89US-00422383
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                                                                                                                                                                                                                                                                          (first entry)
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                                     20; Conservative
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                    Similarity
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                                                                                                                                                                                                                                                                          09-FEB-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                     JS6967029-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-MAY-1995;
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21-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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 Query Match
Best Local 8
                                     Matches
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The invention relates to an aptamer probe (I) comprising an aptamer having an obligonucleotide tail, and a second oligonucleotide having a gequence complementary to at least a portion of a sequence of the oligonucleotide tail comprising an optional label. Also described: (I) detecting (MI) at least one target analyte having at least two binding stites, in a sample; (2) a nanoparticle-aptamer conjugate probe (II) comprising annoparticles, and at least one type of aptamers being present or net nanoparticle at a surface density ranging from between about 1.0x10 10 and about 5.0x10 12 aptamers/cm2; (3) a substrate, at least one type of capture aptamers bound to the substrate, where each type of detecting one or more target analytes comprising a substrate, at least one type of capture aptamers bound to the substrate and arranged in an array of discrete spots, and electrodes located between the discrete spots; and (4) a kit (KI) for detecting one or more analytes in a sample comprising (I) or (III) and an optional substrate. (I) is useful for detecting at least one target analyte in a sample. The method is useful
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disseminated fungus disease, fulminating septicemia, malaria, vitamin B12, and folic acid deficiency, pyridoxine deficiency, Diamond blackfan anemia, hypopigmentation disorders such as piebaldism and vitiligo, and AIDS. (M2) is useful in expanding early hematopoietic progenitors in syngeneic, allogeneic or autologous bone marrow transplantation. (M1) enhances hematopoiesis by expanding early hematopoietic progenitors. The present sequence represents a universal PCR primer for SCF, which is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel aptamer probe comprising aptamer having oligonucleotide tail, and oligonucleotide having sequence complementary to portion of sequence of oligonucleotide tail having optional label, useful for detecting target
                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                     Score 20; DB 1; Length 20;
                                                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                                              Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                               Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic poly-T20 oligonucleotide SEQ ID NO:10.
                                                                                                                                                                                                                                 100.0%; Pred. ...
                                                                                                                                           in an example from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Senical MJ,
                                                                                                                                                                                                                                                                                                     2707 CTAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aptamer; analyte detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                            BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-MAY-2004; 2004US-0567874P, 22-NOV-2004; 2004US-00995051.
                                                                                                                                                                                                                         0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          AEF05127 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-MAR-2006 (first entry)
                                                                                                                                                                                                                   Query Match 0.7
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2006-099415/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            analyte in sample.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEF05127;
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                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 656
                                                                                                                                                                                                                                                                                                                                                                                                                       AEF05127
    88888888888888
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The invention relates to an aptamer probe (I) comprising an aptamer having an oligonucleotide tail, and a second oligonucleotide having a sequence complementary to at least a portion of a sequence of the coligonucleotide tail comprising an optional label. Also described: (I) detecting (MI) at least one target analyte having at least two binding of sites, in a sample; (2) a nanoparticle-aptamer conjugate probe (II) comprising nanoparticles, and at least one type of aptamers being present on the nanoparticle at a surface density ranging from between about on the nanoparticle at a surface density ranging from between about on the nanoparticle at analytes comprising a substrate, at least one type of capture aptamers bound to the substrate, where each type of capture aptamers bound to the substrate, where each type of capture aptamers bonds to the specific target analytes and arranged in an array of discrete spots, and electrodes located between the discrete spots; and (4) a kit (KI) for detecting one or more analytes in a sample of comprising (I) or (II) and an optional substrate. (I) is useful for detecting one or more target analyte in a sample. The with or comprising at least one target analyte in a sample. The kit or substrate are useful in
    is useful in
substrate are useful for detecting one or more analytes. (I) is useful ir barcode detection assays and in therapeutic, diagnostic and target validation applications. (I) can be readily synthesized, manipulated and stored for long periods of time. (I) provides reproducibility in production. The present sequence represents a synthetic poly-T20 oligonucleotide, which is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel aptamer probe comprising aptamer having oligonucleotide tail, and oligonucleotide having sequence complementary to portion of sequence of oligonucleotide tail having optional label, useful for detecting target
                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                          DB 1; Length 20;
6.8e+02;
                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Senical MJ, Garimella V;
                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic poly-A20 oligonucleotide SEQ ID NO:7.
                                                                                                                                                                                                        Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 6.8 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; SEQ ID NO 7; 47pp; English.
                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                 20 AAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                        AEF05124 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-MAY-2004; 2004US-0567874P.
22-NOV-2004; 2004US-00995051.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       aptamer; analyte detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NANO-) NANOSPHERE INC
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                                                                                                                             Invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEF05124;
                                                                                                                                                                                                                                                                                                                                                                                                 657
                                                                                                                                                                                                                                                                                                                                                                                               RESULT 6
AEF05124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rр
                  and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mot\overline{e}_{\rm B} "Phosphorothioate backbone with 2'-O-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotide; phosphorothioate; 2'-0-methoxyethyl; 2'-MOE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Preparing an internucleotide phosphorothioate linkage enriched in the or Sp enantiomer comprises coupling the synthon to the 2'-substituted nucleoside in the presence of the coupling agent.
barcode detection assays and in therapeutic, diagnostic and target validation applications. (1) can be readily synthesized, manipulated stored for long periods of time. (1) provides reproducibility in production. The present sequence represents a synthetic poly-A20 eligonucleotide, which is used in the exemplification of the present
                                                                                                                                                                              Gaps
                                                                                                                                                                              ;
                                                                                                                                                Length 20;
                                                                                                                                                                              Indels
                                                                                                              Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                              0.7%; Score 20; DB 1; L(100.0%; Pred. No. 6.8e+02;
                                                                                                                                                          100.0%; Preα. ....
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide, SEQ ID NO: 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                               2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUN-2001; 2001US-00881535.
02-SEP-2004; 2004US-00932630.
                                                                                                                                                                                                                                                                                                                            AEG11130 standard; DNA; 20
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                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      *tag=
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                                                                                                                                                                Similarity
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modified_base
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Unidentified.
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                                                                                                                                                                                20;
                                                                                       invention.
                                                                                                                                                                                                                                                                                                                                                              AEG11130;
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                                                                                                                                                                  Best Local
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- by amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                                                                                                                                                   A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                       Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
     Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                        TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                               alysis of cDNA and gene expression - digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2706 ACTAAAAAAAAAAAAAA 2725
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                                                                                                                              JP06303997-A.
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                                                                                                                                                                01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                               Analysis of
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                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverses transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                              Gaps
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oligonucleotide with internucleotide phosphorothioate linkage
                                                                                                                                                                                                                                                                                                                                                                               Reverse transcription primer used in cDNA analysis technique.
                                                                     ch 0.7%; Score 20; DB 1; Length 20; 1 Similarity 100.0%; Pred. No. 6.88+02; 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
ive 0; Mismatches 0; Indels
                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          by digestion with restriction enzymes
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                                                                                                                                           2709 AAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                                          04-AUG-1995 (first entry)
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                                                                                       Local Similarity
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nes 20; Conserv
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Matches
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AAQ75713/
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The sequences represented by, AAQ90390-90401 are synthetic DNA probes containing 3' ribonucleoside termini. The sequences shown in AAQ90402-15 are synthetic DNA probes with 5' amino termini. These sequences were specific for the polymorphisms of HLA gene dQa. The sequences were in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and sensitivity of these reactions are greatly improved at micro-locations
                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of ababiled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded CDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New self-addressable electronic devices - used for multi-step and multiplex reactions such as DNA hybridisation(s), clinical diagnostics and bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CP-1; HLA; dQa; 3' ribonucleoside; self-addressable electronic device; SAED; hybridisation; ss.
                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                      0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/note= "3' ribonucleoside terminal"
                                                                                                                                                                                                                      Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                 Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                  2706 ACTAAAAAAAAAAAAAAAA 2725
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                                                                                                                                                                                                                                                                                                                                                       20 ACTAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ90391 standard; DNA; 21 BP.
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                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tu E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-OCT-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ90391;
                                                                                                                                                                                                                                                            Query Match
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The sequences given in AAT10742-67 are synthetic oligonucleotides which are used in the construction of the electronically self-addressable device (BD) of the invention. The BC comprises a substrate, an electrode or opt. a number of electrode supported by the substrate, a current source operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding to the electrode. The attachment layer is capable of attaching a macromolecule. The ED is used for genetic typing and comprises a number of electrodically, such as one of these probes, attached to each of the locations capable of a genetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Electronically self-addressable device - used for electronic control of, e.g. nucleic acid hybridisation.
                                                                                                                                                                                                                                                                                                 Electronically self-addressable device; ED; electrode; current source; attachment layer; permeable; counterion; genetic typing; probe; detection; ss.
                                                                 Gapa
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                                    Length 21;
                                                                 Indels
                           DB 1; Le...
                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "3'-ribonucleoside terminus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
           Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                     0.7%; Score 20; DB
100.0%; Pred. No. 7e+
iive 0; Mismatches
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                                                                                            2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                      1 AAAAAAAAAAAAAAA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94US-00271882
                                                                                                                                                                                                                                                                              Oligonucleotide probe, CP-1
                                                                                                                                                                                            AAT10743 standard; RNA; 21
                                                                                                                                                                                                                                                    (first entry)
                                                                  Conservative
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                                                    Local Similarity
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                       Query Match
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                                                                  Matches
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0.7%; Score 20; DB 1; Length 21; 100.0%; Pred. No. 7e+02; tive 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAAAAA 2728

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Query Match Best Local Similarity

## AAAAAAAAAAAAAA 20

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AAX81302 standard; DNA; 21 BP AAX81302; RESULT 664 AAX81302

20-AUG-1999 (first entry)

Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss. 3' ribonucleoside oligonucleotide probe CP-1.

Synthetic

Location/Qualifiers /\*tag= misc\_RNA

409929711-A1

17-JUN-1999

98WO-US025475. 01-DEC-1998;

97US-00986065. 05-DEC-1997;

(NANO-) NANOGEN INC.

WPI; 1999-385567/32.

Edman CF; Nerenberg MI, Heller MJ, Tu Ε, Butler WF, Sosnowski RG,

New microelectronic device designed to carry out and control multi-step and multiplex molecular biological reactions in microscopic format.

Example 1; Page 89; 179pp; English.

The specification describes a self-addressable, self-assembling microelectronic device which is designed to actively carry out and control multi-step and multiplex molecular biological reactions in microscopic formats. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation layer which overlies the electrode. This permeation layer which overlies the electrode. This permeation layer separates the attached oligonucleotides and hybridization but also separates the attached oligonucleotides and hybridize drarget DNA sequences from the highly reactive electrochemical environment generated immediately at the electrode surface. The microelectronic device is designed and fabricated to activaly carry out and control reactions such as nucleic acid hybridizations, antibody/antigen reactions, sample preparation, diagnostics and biopolymer synthesis. The device can electronically control the transport and attachment of specific binding electronically so nucleic acids and polypeptides, to specific microlocations. The device can subsequently control the transport and reactions of the analytes or reactants at the addressed specific microlocations. The present sequence represents a probe used to exemplify the invention hybridization reactions and improve the detection of analytes. The device is able to concentrate analytes and reactants, remove non-specifically bound molecules, provide stringency control for DNA 

Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;

ö Score 20; DB 1; Length 21; Pred. No. 7e+02; 0; Mismatches 0; Indels Query Match 0.7%; Sco Best Local Similarity 100.0%; Pr Matches 20; Conservative 0; Best Loca Matches

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ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable Rat DNA microarray capture oligonucleotide #7. Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH. ADK01287 standard; DNA; 21 BP. 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. (first entry) Boekenkamp D, Dieck HT, WPI; 2003-714082/68. ss; hybridisation; DE10208794-A1. 04-SEP-2003. 06-MAY-2004 Rattus sp. ADK01287; RESULT 66 ADK01287/ 

Example; Page 5; 8pp; German. and constant regions.

This interior describes a novel method for sorting single-stranded concluded acids by isolation and hybridisation of nucleic acids pools, then reading out, where the nucleic acids are selectively bound using capture cadening out, where the nucleic acids are selectively bound using capture cadening out, where the nucleic acids are selectively bound using capture cadening out, where the nucleic acids are surface of a solid matrix and (b) comprise variable anchor region, the complement of which is present at least nucleic acid and a 3' variable, discriminatory region that comprises all possible combinations of up to 10 nuclectides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) conducting to the surface is (quasi) conducting to the surface acid in for analysis of patterns, especially in mucosal, conducting to a carboxylic or fatty acid) or through a molecular, conductives or supplements, especially minerals, trace elements, organic acids in a callows inaging of the complete pattern of all nucleic acids from cells, and conjucte pattern of all nucleic acid pool. Since the mathod is based on comparison of nucleic acid pools of the invarious nucleic acid in the mathod is passible. AbkOl1344 represent converse and in the nucleic acid pools. This invention describes a novel method for sorting single-stranded capture probes used in the method of the invention.

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

0.7%; Score 20; DB 1; Length 21; 00.0%; Pred. No. 7e+02; ve 0; Mismatches 0; Indels 100.0%; Pred. No. ... Query Match Best Local Similarity 100.0 Matches 20; Conservative

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2707 CTAAAAAAAAAAAAAAA 2726
          CTAAAAAAAAAAAAAAAA 1
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ADK01285/c ID ADK01285 standard; DNA; 21 BP. 셤

ADK01285;

Rat DNA microarray capture oligonucleotide #5.

(first entry)

06-MAY-2004

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Dieck HT, Hoppe H; Boekenkamp D,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable WPI; 2003-714082/68

Example; Page 4; 8pp; German.

and constant regions

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable and non-variable regions. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture of particularly locked nucleic acids (INA) and the anchor region comprises are particularly locked nucleic acids (INA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streaptavidin. The metrial and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, callines or supplements, especially minerals, trace elements, organic called sandour nutritional compounds, e.g. food activity of pharmaccuticals and/or nutritional compounds, e.g. food activity of pharmaccuticals and/or nutritional compounds, e.g. food activity of analysis of patterns, especially minerals, trace elements, organic callows inaging of the complete pattern of all nucleic acid so and electricate and electricate and electricate and activity of analysis of the complete pattern of all nucleic acid so and electricate and electricate and acid pools of nucleic acid pool. Since the captes matrix minaturisation is possible, how acid pools in a particular pattern of allows matrix minaturisation is possible, how acid pools and particular pattern of t capture probes used in the method of the invention.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using and (b) agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture of agents are particularly locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface interaction with streptavidin. The metrial and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) molecular filaments and binding of shingle conducting propertical field or through a molecular situated by an electrical field or through a molecular of physical, stimulated by an electrical field or through a molecular conducting propertical sud/or nutritional compounds, e.g. food activity of pharmaccuticals and/or nutritional compounds, e.g. food activity of pharmaccuticals and/or nutritional compounds, e.g. food activity of analysis of patterns, specially minerals, trace elements, organic caids indino, carboxylic or fatty acid) or their derivatives, salts and contrast maining of the complete pattern of all nucleic acid singing of the complete pattern of all nucleic acid sold in a cell, and can lows imaging of the complete pattern of all nucleic acid sold on the nucleic acid pools, not individual can detect very small differences in the nucleic acid pools, not individual can detect very small of the method is ba

Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;

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                                                                                                                                                         capture oligonucleotide, pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                  Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                 Gaps
                 ö
Length 21;
0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                          Rat DNA microarray capture oligonucleotide #6.
                                  2707 CTAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                    Hoppe H;
                                                                                                                                                                                                                                                                     (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                  CTAAAAAAAAAAAAAAAA 1
                                                                                          BP.
                                                                                                                                                                                                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                          ADK01286 standard; DNA; 21
                                                                                                                          (first entry)
  Ouery Match
Best Local Similarity 100.8
Matches 20; Conservative
                                                                                                                                                                                                                                                                                     Boekenkamp D, Dieck HT,
                                                                                                                                                                                                                                                                                                                                     and constant regions.
                                                                                                                                                                                                                                                                                                     WPI; 2003-714082/68
                                                                                                                                                          hybridisation;
od; nerve; germ
                                                                                                                                                                                                    DE10208794-A1.
                                                                                                                                                            ss; hybridisat
blood; nerve;
                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                   04-SEP-2003.
                                                                                                                                                                                    Rattus sp.
                                                                                                           ADK01286;
                                                  20
                                                                           RESULT 667
                                                                                    ADK01286/
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture of reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture agents are particularly locked nucleic acids. The capture of particularly locked nucleic acids (IMA) and the annor region comprises as sequence of 10-50, particularly 15-25, T residues. The acpture of interaction with streptavidin. The matrix is of plastic, ceramic, glass, cracial, reain, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of single stranded nucleic acids to the surface is quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional capturity of micro acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                    Gaps
                                                                                                                                      ö
                                                                      DB 1; Le...
  capture probes used in the method of the invention.
                                         Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rat DNA microarray capture oligonucleotide #63.
                                                                                    0.7%; Score 20; DB
100.0%; Pred. No. 7e+
ive 0; Mismatches
                                                                                                                                                                        2707 CTAAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Boekenkamp D, Dieck HT, Hoppe H;
                                                                                                                                                                                                    20 CTAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                            ADK01343 standard; DNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                               20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-714082/68
                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                          ADK01343;
                                                                                      Query Match
                                                                                                                               Matches
                                                                                                                                                                                                                                                                                   RESULT 66
SXS
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides of compariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of single stranded nucleic acids. The capture comparises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptaviain The matrix is of plastic, ceramic, glass, capture oligonucleotides are biotinylated and immobilised on a surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, attinity of pharmaceuticals and/or nutritional compounds, e.g. food
                                                                                                                                                                                                               ö
      and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                               ö
                                                                                                                                                                   0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                          Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat DNA microarray capture oligonucleotide #51.
                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hoppe H;
                                                                                                                                                                                                                                                                                            20 AAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                  ADK01331 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                 Query Match 0.7
Best Local Similarity 100.
Matches 20, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boekenkamp D, Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK01331;
                                                                                                                                                                                                                                                                                                                                                           RESULT 669
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and acide (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and repreducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, an method is based on comparison of nucleic acid pool. Since the method is based on comparison in possible, ADK01281-ADK01344 represent capture probes used in the method of the invention. additives or supplements, especially minerals, trace elements, organic 88888888888888

Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö DB 1; Le... 0.7%; Scor. V 100.0%; Pred. No. .. '... 0; Mismatches 2708 TAAAAAAAAAAAAAAA 2727 20 TAAAAAAAAAAAAAAA 1 Local Similarity 100. Query Match Matches ò 셤

ВР ADK01329 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01329 RESULT 670 ADK01329, 

Rat DNA microarray capture oligonucleotide #49.

98; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions. Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acids pools, then reading out, where the nucleic acids are selectively bound using capture capture the nucleic acids are selectively bound using capture is membhilised on the surface of a solid marrix and (b) comprise variable anchor region. The capture oligonucleotides that capture as 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular,

physical, stimulated by an electrical field or through a molecular sieve The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, an can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. 

Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;

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ö Gaps ö 0.7%; Score 20; DB 1; Length 21; 00.0%; Pred. No. 7e+02; ve 0; Mismatches 0; Indels 100.0%; Pred. worlder 20; Conservative Query Match Best Local Similarity Matches

2708 TAAAAAAAAAAAAAAAA 2727 

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ADK01332 standard; DNA; 21 06-MAY-2004 ADK01332; ADK01332/ RESULT 

ВЪ.

(first entry)

Rat DNA microarray capture oligonucleotide #52.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT,

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WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particulary locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass,

metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, the method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acid from cells. It can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturiation is possible, ADK01281-ADK01344 represent capture probes used in the method of the invention.

Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

.; 0 0.7%; Score 20; DB 1; Length 21; 100.0%; Pred. No. 7e+02; iive 0; Mismatches 0; Indels 2708 TAAAAAAAAAAAAAAA 2727 20 TAAAAAAAAAAAAAA 1 Local Similarity 100. 1es 20; Conservative Query Match Matches ઠે g

ADK01342 standard; DNA; 21 BP (first entry) 06-MAY-2004 ADK01342; RESULT 672 ADK01342/c 

Rat DNA microarray capture oligonucleotide #62.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5' invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture

comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical filed or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the cerivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible crepresentation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid in a cell, and method is based on comparison of mucleic acid pool. Since the method is based on comparison of nucleic acid pools individual capture probes used in the method of the invention.

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Sequence 21 BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;

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Gaps

Gaps ö Score 20; DB 1; Length 21; Pred. No. 7e+02; 0; Mismatches 0; Indels 100.0%; Pred. ... 0.78; 20; Conservative Similarity Query Match Best Local & Matches

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2709 AAAAAAAAAAAAAAA 2728 AAAAAAAAAAAAAAAAAA 20

ઠે 셤 RESULT 673 ABD25908

ВР ABD25908 standard; DNA; 21

ABD25908;

(first entry) 29-JUL-2004

AI654215-derived oligonucleotide SEQ ID 4920.

Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory districtses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

THE STATE OF STATE OF

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka , Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

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The invention describes a nover composition, alleviating active agent, comportating adenosine (effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating or conference in apparate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cycostatic activity, is a certain a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated inflammation, allergies, asthma, impeded respiration; respiratory cy distress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary charteress syndrome, pain, cystic fibrosis, allergic thinitis, pulmonary cor thymidines present in the target RNA serves to prevent the bystem or the oligonucleotides into products that free adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine to prevent the system of the oligonucleotides into products that free adenosine content of the oligonucleotides into products that free adenosine or prevent and the present in the reduce the productio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                          This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AI654215-derived oligonucleotide SEQ ID 4919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                   Claim 15; SEQ ID NO 4920; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAAAA 2729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AAAAAAAAAAAAAAAAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD25907 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 95.2'
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Katz E, Claim 15; SEQ ID NO 4919; 763pp; English. brain, heart, kidney, etc, unwanted effects due to it 2709 AAAAAAAAAAAAAAAAAA 2729 Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 1 AAAAAAAAAAAAAAAAAAAAAAAAAA 21 BP. (EPIG-) EPIGENESIS PHARM INC. 95.2%; ADK67451 standard; DNA; 21 06-MAY-2004 (first entry) Best Local Similarity 95.2 Matches 20; Conservative bronchodilating agent. WPI; 2003-093058/08. prevent any Miller S, ADK67451; Query Match Nyce JW, RESULT 675 ADK67451 g ð ö Gaps ö 0.7%; Score 20; DB 1; Length 21; 95.2%; Pred. No. 7e+02; 1; Indels

Thus invention describes a novel composition, a latter active agent, comparising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and content adequate associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or mallignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung area ö inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to invention describes a novel composition (a) a first active agent Gape Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and ä ö Aguilar 0.7%; Score 20; DB 1; Length 21; 15.2%; Pred. No. 7e+02; 1; Indels intercalator; electrochemical detection; mismatch; ds. Electrochemical detection intercalator-related DNA 1. Pabalan J, Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 1 Other; 0; Mismatches JP2004024114-A. Synthetic  335

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RESULT 677
                                                                                                         Matches
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                                                                                                                                      The intercalator of the invention may be useful for the electrochemical detection of a gene, as an electrochemical double stranded DNA detection reagent and as an intercalator for inhibiting the influence of mismatch DNA and single stranded DNA. The intercalator enables the transmission of electronic transition between two base pairs to occur efficiently. The current sequence is that of the electrochemical detection intercalator related DNA 1 of the invention.
                                                                                                                              The invention relates to a novel intercalator having a specific formula.
                                                                                                                                                                                                                                         Gaps
                                                                                      Novel intercalator, useful as electrochemical double stranded DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel fluorescently labeled nucleoside triphosphate, useful for
                                                                                                                                                                                                                       0.7%; Score 20; DB 1; Length 21;
100.0%; Pred. No. 7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                      Sequence 21 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                             RNA, SEQ ID NO: 6 used in sequencing nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                              'note= "Biotinylated"
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                                                                                                              Example 1; Page 23; 24pp; Japanese.
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AEB80248 standard; RNA; 21
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                                               TAKENAKA S.
TUM KENKYUSHO
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es 20; Conserv
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                                                                                               detection reagent
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(BUZB/) BUZBY P
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29-JAN-2004
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                                               (TAKE/)
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multiple amino
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                                                                                   The present invention relates to fluorescently labeled nucleoside triphosphate. The invention is useful for determining nucleic acid sequence of target nucleic acid. The present sequence is a RNA used in sequencing nucleic acids.
                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amino modified oligodeoxyribonucleotide; oligonucleotide; achiral linker reagent; 5-(aminomethyl)-1,3-benzenedimethanol; N-fluresceinyl-(5-aminomethyl)-1,3-benzenedimethanol; hybridisation probe; PCR primer; nucleic acid sequencing; affinity matrix; cloning recombinant DNA; in-vitro mutagenesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New achiral linker reagents - useful for incorporation of gps. or reporter gps. into oligo:nucleotide(s).
determining nucleic acid sequence of target nucleic acid
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100.0%; Pred. No. 7e+
ive 0; Mismatches
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                                         SEQ ID NO 6; 39pp; English
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EGHOLM M.
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                                                                                                                                                                                                                                                              Similarity
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                                         Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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(EGHO/)
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(DAHL/)
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or reporter groups into oligonucleotides. They are compatible with conventional DNA synthesis following the phosphoramidite methodology, and can be incorporated in good yields. The linker reagents may be used for labelling of oligonucleotides. They may also be used for preparation of oligonucleotides. They may also be used for preparation of primers in the polymerase chain reaction or in nucleic acid sequencing reactions, for production of affinity matrices for purification of DNA binding proceins or other biomolecules, for production of affinity matrices for detection of nucleic acid sequences, for cloning recombinant
                                                                                                                                                                                                                                                                                         DNA or for in-vitro mutagenesis
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Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;

ö Query Match

0.7%; Score 20; DB 1; Length 22;
Best Local Similarity 90.9%; Pred. No. 7.2e+02;
Matches 20; Conservative 0; Mismatches 2; Indels 2709 AAAAAAAAAAAAAAAAAA 2730 ò

22 AAAAAAAANNAAAAAAAA 1 유

Oligo dT primer #4 used in method to study gene expression. Oligo dT primer; gene expression analysis; primer; ss ABK86172 standard; DNA; 24 (first entry) WO200236828-A2. 24-SEP-2002 Synthetic. ABK86172; RESULT 678 

01-NOV-2001; 2001WO-US045401. 01-NOV-2000; 2000US-0244933P.

10-MAY-2002

Dombkowski AA, Nagel AC; Kane MD,

(GENO-) GENOMIC SOLUTIONS INC.

WPI; 2002-508123/54.

Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identimer having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.

Example 1; Page 15; 45pp; English.

of gene expression in one or more samples, comprising an identification of gene expression in one or more samples, comprising an identification of proceed as a specific oligo-dT primer sequence, where the identifier a variety of detectable marker at its 5' end. The system is useful for identifying any or all genes expressed in a given in vivo or in vitro RNA sample, as well as the relative differences in mRNA between 2 or more samples, where that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA capturession. The system has the following advantages over existing expression. The system has the following advantages over existing nethods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence inferences information concerning changes or differences The invention relates to systems for identification and characterisation

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in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates cDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology techniques; and (d) does not require prior knowledge of the sequence of the genome of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present sequence represents an oligo dT primer used in the method of the
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                                                                                                                                                                                                                                                   Length 24;
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100.0%; Pred. No. 7.5e+02;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                Conservative
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2709 AAAAAAAAAAAAAAAAA 2728 24 AAAAAAAAAAAAAA S ò 셤

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Gaps

ВР 24 (first entry) ADG75987 standard; DNA; 11-MAR-2004 ADG75987; RESULT 679 ADG75987/c 

Immunostimulatory non-CpG oligonucleotide IMT 059 SeqID 98.

ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma

Synthetic.

WO2003101375-A2

11-DEC-2003.

30-MAY-2003; 2003WO-EP005691.

30-MAY-2002; 2002CA-02388049

(IMMU-) IMMUNOTECH SA.

Lopez RA;

WPI; 2004-053333/05.

New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. melanoma

Disclosure, Fig 3; 139pp; English.

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a cpG motif), which can atimulate an immune response in animals of the order of primate, including humans. The immune and antibody production in B-cells, as well as cell differentiation, cytokine and antibody production in B-cells, as well as cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG

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Sequence 25 BP; 21 A; 1 C; 2 G; 1 T; 0 U; 0 Other;
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     is used in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                           Hypermutable organism; dominant negative allele; mismatch repair gene; spontaneous mutation; DNA repair; purine nucleotide phosphorylase; PNP;
                                                                                     Gaps
variant DNA oligo, used in an exemplification of the invention.
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                                                         Length 24;
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                           Sequence 24 BP; 0 A; 0 C; 4 G; 20 T; 0 U; 0 Other;
                                                      0.7%; Score 20; DB 1; Le
100.0%; Pred. No. 7.5e+02;
tive 0; Mismatches 0;
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6. 25
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/note= "In c
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"In-frame polyA tract"
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                                                                                                                                                                                                             AAD26900 standard; DNA; 25 BP.
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                                                                                                                                                                                                                                                                     09-APR-2002 (first entry)
                                                                                20; Conservative
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NICOLAIDES N C.
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                                                                    Sest Local Similarity
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GRASSO L.
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                                                                                                                                                                                                                                                                                                                                                                                 Bacteria.
Unidentified.
Chimeric.
                                                                                                                                                                                                                                                                                                                                                        bacteria; 88.
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(SASS/)
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(VOGE/)
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                                                                                Matches
                                                                                                                                                                                 RESULT
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The invention relates to a method of determining a functional property of a fluid in a chamber comprising introducing into the chamber a support to which is bound several test elements each of the test elements comprising a reaction domain and a detection domain, introducing into the chamber a fluid that is interactive with the reaction domains, removing the fluid from the chamber, determining by means of the detection domains the fluid as not interacted with the reaction domains the locations at which the fluid has not interacted with the reaction domains comprises a member of a specific binding pair. The determining of the step involves treating the test elements to modify only those reaction comprises a member of a specific binding pair. The determining of the step involves treating the test elements to modify only those reaction comprises that have interacted with the fluid. The fluid in the method is chosen from the flow pattern of the fluid. The fluid in a chamber and the method provides for the characterisation of a fluid in a known fluid in a new flow cell or a new fluid in a known flow the invention. This sequence represents a test element used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Determining a functional property of fluid in chamber by introducing a support comprising test elements having reaction and detection domains, introducing a test fluid, and detecting locations not reactive with the
                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fluid functional property; fluid flow pattern;
fluid reagent distribution; time dependent fluid reactivity; ss.
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0.7%; Score 20; DB 1; Length 25; 100.0%; Pred. No. 7.6e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                         2709 AAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                6 AAAAAAAAAAAAAAA 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Test element oligonucleotide #1
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                                                                                                                                                                                                                                                                                                                                                                                               ADH78589 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Leproust EM, Amorese DA,
                                 1 Similarity 100.
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (LEPR/) LEPROUST E M. (AMOR/) AMORESE D A. (PECK/) PECK B J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-061269/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
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Query Match
Best Local S
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The invention relates to an oligonucleotide having the formula: (a) 5'

XYMIYZN2 3' where 5' denotes the 5' end of the oligonucleotide and 3'
denotes the 3' end of the oligonucleotide, x is a T or modified T
cucleotide, y is a C or modified C nucleotide, z is a G or modified G
nucleotide, and N1 and N2 are polymucleotides that do not include a CG
dinucleotide, where N1 does not include 5' Z nucleotide, and where a 3'
concleotide consisting of the YZ dinucleotide and the N2
polymucleotide consisting of the YZ dinucleotide and the N2
polymucleotide contains a number of nucleotides that is at most 45% of
the number of nucleotides in the oligonucleotide, and (b) 5' XYNIYZNZ 3'
where 5' denotes the 5' end of the oligonucleotide, and (b) 5' XYNIYZNZ 3'
where 5' denotes the 5' end of the oligonucleotide, and (b) 5' XYNIYZNZ 3'
c not of the oligonucleotide, where X is a T or modified T nucleotide, N is a
c or modified c nucleotide, where N1 does not include a CG
dinucleotide of 5 to 3 nucleotides. Also described are: (1) a
polymucleotide of 5 to 3 nucleotides. Also described are: (1) a
polymucleotide of 5 to 3 nucleotides. Also described are: (1)
a polymucleotide of 5 to 3 nucleotides. Also described are: (1)
c apparentic agents, monoclonal antibodies, and anticancer agents,
radiotherapeutic agents, monoclonal antibodies, and anticancer agents,
c netto of interferon (IFN)-alpha expression relative to IFN-alpha
administering an oligonucleotide or a pharmaceutical composition to the
subject in need of increased IL-10 expression relative to IFN-alpha
cyll response in a subject exposed to an antigen; (4) a method of
inducing an antigen-specific regulatory B cell response in a subject by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus.
                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                        pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.
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0
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     DB 1; Length 25; 7.6e+02;
                                                  0; Indels
Query Match 0.7%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 7.6 Matches 20; Conservative 0; Mismatches
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                                                                                                 2707 CTAAAAAAAAAAAAAAAA 2726
                                                                                                                                                 CTAAAAAAAAAAAAAA 25
                                                                                                                                                                                                                                                                       AED81293 standard; DNA; 23 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                      AED81293;
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administering an immunostimulatory nucleic acid or composition to a subject exposed to an antigen; (5) a method of treating an allergy or aethma by exposing a subject to an allergen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or allergiate an allergen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or treat an autoimmune disease in the subject, and (7) a method of reducing an antigen-specific response to an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigen-specific response to the implant in the subject. The oligonucleotide and antigen-specific response to the implant in the subject. The oligonucleotide incompositions of the invention are useful for treating allergies, asthma, autoimmune of the invention are useful for treating allergies, asthma, autoimmune composition are useful for treating allergies, asthma, autoimmune neuropathies such as Gaillain-Barre, autoimmune thrombocypenia, temporal arterties, systemic lupus erythematosus, multiple sclerosis, temporal arterties, anti-phospholipid syndrome, psoriasis, pemphigus vingaris, vasculties such as Wegener's granulomatosis, utcoimmune compositions of temporal arterties, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculties and as Wegener's granulomatosis, utcoimmune composities and as Wegener's granulomatosis, utcoimmune mediated diabetes mellitus, grave's biesase, leading allergies autoimmune composities and as a parkylosities, autoimmune composities and as a paganese of the adrend glabetes mellitus, grave's biesase, leading allergies, autoimmune comp polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present 

Sequence 23 BP; 0 A; 1 C; 1 G; 21 T; 0 U; 0 Other;

Gaps ö Length 23; 2; Indels DB 1; Score 19.8; DB 1; Pred. No. 7.5e+02; 0; Mismatches 2 0.7%; Local Similarity 91.3 nes 21; Conservative Query Match Matches

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2709 AAAAAAAAAAAAAAAAAAAAAAAA 2731 23 AAAAAAAAAACGAAAAAAA 1 ð 셤

Human FD 17 PCR primer 2 SEQ ID NO:4. ВЪ AAH44623 standard; DNA; 24 (first entry) 16-NOV-2001 AAH44623; RESULT 683 AAH44623/c

Human, FD 17; cytostatic; virucidal; immunomodulatory; haemostatic; antiinflammatory; gene therapy; malignant tumour; haemopathy; human immunodeficiency virus infection; HIV infection; immunological disease; inflammation; PCR primer; 88. 26-FEB-2001; 2001WO-CN000221. 02-MAR-2000; 2000CN-00111868. WO200164729-A1. Homo sapiens. 07-SEP-2001. 

(BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.

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Gaps

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The present invention describes the human FD 17 protein (I). (I) has extostatic, virucidal, immunomodulatory, antiinflammatory and haemostatic activities. The polynucleotide encoding (I) can be used in gene therapy. (I) and the polynucleotide encoding it are applicable in the diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents a PCR primer for human FD 17, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide-human macroprotein 21.78 and polynucleotide encoding it, for treating diseases such as embryo development teratogenesis and tumor.
                                                                                  New human polypeptide FD 17 for diagnosing and treating malignant tumor, hemopathy, human immunodeficiency virus (HIV) infection, immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes human macroprotein 21.78 (I). Also described is a process for preparing (I) using DNA recombination techniques. (I) and the polynucleotide sequence encoding it (II) can be used in the treatment of diseases such as embryo development teratogenesis and tumours. The present sequence represents a PCR primer for (I), which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; macroprotein 21.78; embryo development teratogenesis; tumour;
PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                    Sequence 24 BP; 0 A; 2 C; 1 G; 21 T; 0 U; 0 Other;
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Pred. No. 7.7e+02;
0; Mismatches 2
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                                                                                                                                                         Example 2; Page 11; 36pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.7%;
Best Local Similarity 91.3%;
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                                                                                                                       diseases and inflammations.
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                                                   WPI; 2001-550164/61.
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                  Mao Y,
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gene expression - by amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate
                                                                                                                                                                                                                                                                                         Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
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    Length 24;
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                               2; Indels
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0.7%; Score 19.8; DB 1;
ilarity 91.3%; Pred. No. 7.7e+02;
Conservative 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        alysis of cDNA and gene expression - digestion with restriction enzymes.
                                                            2709 AAAAAAAAAAAAAAAAAAA 2731
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AAQ75648 standard; DNA; 21
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               Similarity
Query Match
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Matches 21; (
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                                                                                                                                                                                                                                                                                                                                         Synthetic.
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WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                                                                   A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                          0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                         Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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 aggregate; restriction enzyme; ss.
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Matches 20; Conserv
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type clabelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                                                                                      Score 19.4; DB 1; Length 21;
Pred. No. 7.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
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Best Local Similarity 95.2%;
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les 20; Conservative
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                                                               Reverse transcription primer used in cDNA analysis technique.
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Pred. No. 7.6e+02;
0; Mismatches 1; Indels
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         AAQ75643 standard; DNA; 21
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AAQ75643/c
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEC files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                      ilysis of cDNA and gene expression - by amplification of mRNA followed digestion with restriction enzymes.
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restriction enzyme; ss.
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95.2%; Pred. No. 7.6e+02;
Ive 0; Mismatches 1; Indels
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& TELEPHONE CORP.
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(NITE ) NIPPON TELEGRAPH
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nes 20; Conserv
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         Length 21;
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       0.7%; Score 19.4; DB 1;
ilarity 95.2%; Pred. No. 7.6e+02;
Conservative 0; Mismatches 1;
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                                                            AAQ75753 standard; DNA; 21 BP.
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ilarity 95.2%;
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Best Local Similarity
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A method for the analysis of cDNA comprises (a) preparing an aggregate of
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Pred. No. 7.6e+02;
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 7.6e+02;
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aggregate; restriction enzyme; ss.
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RESULT 695 AAQ75716/

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                         digestion with restriction enzymes.
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0.7%; Score 19.4;

Query Match

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95.2%; Pred. No. 7.6e+02;
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Matches 20; Conservative
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                                                                                                                           AAQ75708,
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labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)
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                                                                                                                                                                                                                                                                                                                           aggregate;
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ID AAQ75
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                                                                                                                                                      gene expression - by amplification of mRNA followed
                                                                                                                                                                                                         A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mRNA followed
                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification of
                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                          (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
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                                                                                                                                                    Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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Best Local Similarity 95.2.
Best Local Similarity
Local Similarity
Local Similarity
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and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                 Query Match 0.7%; Score 19.4; DB 1; Length 21; Best Local Similarity 95.2%; Pred. No. 7.6e+02; Matches 20; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                    Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        restriction enzyme; ss
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AAQ75696 standard; DNA;
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                                                                  Reverse transcription primer used in cDNA analysis technique.
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ilarity 95.2%; Pred. No. 7.6e+02;
Conservative 0; Mismatches 1: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                    by digestion with restriction enzymes.
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             AAQ75721;
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Matches
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                                               Analysis of cDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypervariable region; ENV protein; vaccinia virus; gag gene; retrovirus; vaccines; infection; protection; primer; ss.
                                                                                                                            A method for the analysis of cDNA comprises (a) preparing an aggregate c double-stranded cDNAs by using an aggregate of mRNAs and a plural type c labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) the double-stranded cDNAs with restriction enzyme and; (c) method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                                         Score 19.4; DB 1; Length 21;
Pred. No. 7.6e+02;
0; Mismatches 1; Indels
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                                                                  by digestion with restriction enzymes.
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                                                                                                     Disclosure; Page 8; 11pp; Japanese.
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              WPI; 1995-018287/03.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Determining complementarity of nucleotide fragment for gene analysis, by comparing flow of electric current from or to electroconductive substrate through DNA fragment, with reference obtained from its complement.
culturing the host cell produces particulate structures containing the fusion gag protein. The recombinant vaccinia virus or the fusion gag protein particles may be used in the production of vaccines for protecting against infection with retroviruses such as HIV
                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                nucleic acid; gene analysis; polymorphism; variation;
                                                                                                                                                                                                                                             Complementary nucleic acid detection method related sequence #5.
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                                                                    Query Match 0.7%; Score 19.4; DB 1; Length 21; Best Local Similarity 95.2%; Pred. No. 7.6e+02; Matches 20; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Takenaka S,
                                                 Sequence 21 BP; 19 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 19.4; DB 1;
95.2%; Pred. No. 7.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     Takagi M,
                                                                                                           2707 CTAAAAAAAAAAAAAAA 2727
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                                                                                                                               CCAAAAAAAAAAAAAAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 12; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                  (FUJF ) FUJI PHOTO FILM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABX79794 standard; cDNA; 21 BP
                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                         07-JUN-2000; 2000EP-00112235.
                                                                                                                                                                                                                                                                                                                                                                              99JP-00159339.
                                                                                                                                                                                                                                                                                                                                                                                                                      Ogawa M,
                                                                                                                                                                                 AAF24290 standard; DNA; 21
                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20; Conservative
                                                                                                                                                                                                                                                                           DNA chip; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-140003/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     Makino Y, Abe Y,
                                                                                                                                                                                                                                                                 Complementary
                                                                                                                                                                                                                                                                                                                   EP1065278-A2
                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1999;
                                                                                                                                                                                                                          03-APR-2001
                                                                                                                                                                                                                                                                                                                                      03-JAN-2001
                                                                                                                                                                                                     AAF24290;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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The invention discloses a method for identifying a candidate polymorphic repeat within a coding sequence (expressed sequence tag, EST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in human genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar atrophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
                                                                                                                                       EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPOUS; Rep-X; human; genetic disease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a candidate polymorphic repeat within a coding sequence, funderstanding or treating genetic disease, comprises detecting tandem repeats in a target coding sequence and scoring the repeats for polymorphic probability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                       EST polymorphic DNA repeat polynucleotide #119.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat DNA microarray capture oligonucleotide #29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fondon JW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Col 495; 588pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAATAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00475947.
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   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TEXA ) UNIV TEXAS SYSTEM.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wren JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-208818/20.
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                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6472154-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-DEC-1999;
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17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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This invention describes a novel method for sorting single-stranded nucleic acide by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceranded nucleic acids to the serial and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of single curanded nucleic acids to the surface is (quasi) covalent, supramolecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (anino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible caid anilows inaging of the complete pattern of all nucleic acid soil an cell, and callows inaging of the complete pattern of all nucleic acid soil and enterior of allierences in the nucleic acid pool. Since the callows inaging matrix anilatery and passible, AbRO1281-AbRO1344 represent converses and an enterior of the method is passible, the method is passed on comparison of nucleic acid pools.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                capture probes used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-714082/68.
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                                                                                                         Rattus sp.
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0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels
                                                                  2705 TACTAAAAAAAAAAAAAAA 2725
                                                                                                21 TACAAAAAAAAAAAAAAA 1
                   Best Local Similarity 95.2
Matches 20; Conservative
    Query Match
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ADK01314 standard; DNA; 21
                                                                        06-MAY-2004 (first entry)
                                     ADK01314;
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RESULT 709 ADK01314,

#XXXX

DNA microarray capture oligonucleotide #34.

capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. blood; nerve; germ hybridisation; 

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Dieck HT, Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acids by isolation are all possible cacids are selectively bound using capture capture that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable respons. The capture oligonucleotides confirmed anchor region, the comprises all possible combinations of up to 10 nucleotides to allow binding of particular sortes of single stranded nucleic acids. The capture capture particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, interaction with streptavidin. The matrix is of plastic, ceramic, glass, comprises and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular steve. The method is used (1) for analysis of patterns, especially in mucosal, captures. The method provides rapid, inexpensive and especially in a cell and active or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and active furty of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic caids (amino, carboxylic or fatty acid) or their derivatives, salts and erget very small differences in the nucleic acids from cells, and dect very small differences in the nucleic acids from cells, and thouse and the compatison of n

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels 20; Conservative Query Match Best Local Similarity Matches

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RESULT 710 ADK01333/

ADK01333 standard; DNA; 21 BP.

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ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                     Rat DNA microarray capture oligonucleotide #53.
                                                                          28-FEB-2002; 2002DE-01008794.
                                                                  28-FEB-2002; 2002DE-01008794
            06-MAY-2004 (first entry)
                                                DE10208794-A1.
                                                         04-SEP-2003.
                                         Rattus sp.
    ADK01333;
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Boekenkamp D, Dieck HT, Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH. WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streaglavidin. The matrix is of plastic, ceramic, glass, ceranded nucleic acids and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, correspondents, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and acityity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible can detect very small differences in the nucleic acid pool. Since the represent or probes used in the method of the invention. capture probes used in the method of the invention

Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;

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Query Match 0.7%; Score 19.4; DB 1; Length 21; Best Local Similarity 95.2%; Pred. No. 7.6e+02; Matches 20; Conservative 0; Mismatches 1; Indels
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ADK01340 standard; DNA; 21 BP. ADK01340/ 

ADK01340;

Rat DNA microarray capture oligonucleotide #60. (first entry) 06-MAY-2004

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nuclectides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are biotinylated and immobilised on a surface by capture oligonucleotides are biotinylated and immobilised on a surface by comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cating famino, carboxylic or fatty acid) or thair derivatives or supplements, especially minerals, trace elements, organic acids and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, salts and cailes and capture and reproducible can lows inaging of the complete pattern of all nucleic acid side the callows inaging of the complete pattern of all nucleic acid pool. Since the callows inaging miniaturisation of nucleic acid pools, not individual comparison of nucleic acid pools, not individual captures matrix and public the method is p capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;

.; 0

Gaps

; 0

Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels 20; Conservative Query Match Best Local Similarity Best Loca Matches

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2709 AAAAAAAAAAAAAAAAAAA 2729
        21
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ADK01284 standard; DNA; 21 BP. (first entry) 06-MAY-2004 ADK01284; 

Rat DNA microarray capture oligonucleotide #4.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattue sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Dieck HT,

Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Sxample, Page 4; 8pp; German.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then nucleic acids by isolation and hybridisation of nucleic acid pools, then capture oligonout where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region. The comprises all possible combinations of up to 10 nuclectides to allow binding of particularly locked nucleic acids. The capture agents are particularly locked nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of shingle stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, surface is (auss) covalent, supramolecular, chair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible caid and allows imaging of the complete pattern of all nucleic acid pool: Since the callows inaging matrix anial differences in the nucleic acid pool: Since the callows matrix miniaturisation is possible. AbX01281-AbX01344 represent and in the method is based on comparison of the invention. capture probes used in the method of the invention.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides consists and comprises variable and non-variable regions. The capture oligonucleotides to have a 5'-invariable anchor region. The complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The metrial and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo) molecular filaments and binding of surface is granded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, catis and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic caids (anino, carboxylic or farty acid) inexpensive and reproducible conjete pattern of allows inaging of the complete pattern of all nucleic acid since the callows inaging of the complete pattern of all nucleic acid acid in a cell, and can lows inaging of the complete pattern of allows in the nucleic acid bools, not individual can detect very small differences in the nucleic acid pool; Since the method is based on comparison of nucleic acid pools, not individual

0 U; 0 Other; Seguence 21 BP; 2 A; 0 C; 0 G; 19 T;

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                                                                                                                                                          capture oligonucleotide, pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                     Gaps
                     ;
0
     Length 21;
    Score 19.4; DB 1; Length 2
Pred. No. 7.6e+02;
); Mismatches 1; Indels
                                                                                                                                            Rat DNA microarray capture oligonucleotide #13.
                                     2706 ACTAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                    Hoppe H;
0.7%; Scur-
95.2%; Pred
                                                     21 ATTAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                   (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                              BP.
                                                                                                                                                                                                                                                   28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                             ADK01293 standard; DNA; 21
                                                                                                                             06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                   Boekenkamp D, Dieck HT,
                     20; Conservative
                                                                                                                                                                                                                                                                                                   WPI; 2003-714082/68
                                                                                                                                                             ss; hybridisation;
blood; nerve; germ
     Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                     DE10208794-A1.
                                                                                                                                                                                                                     04-SEP-2003.
                                                                                                                                                                                      Rattus sp.
                                                                                                              ADK01293;
                                                                               RESULT 713
                                                                                       ADK01293/c
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture of agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides of have a 5. invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the annor region comprises a sequence of 10-50, particularly 15-25, T residues. The agents are particularly lockof nucleic acids (LMA) and the annor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streeptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve or supplements, especially minerals, trace elements, organic cativity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional captavity is alter and mixtures. The method provides rapid, inexpensive and reproducible cacids (amino, carboxyllic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and representation of differences in pools of nucleic acids ecids or a percentals.
                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                               .;
0
                                                                                                                                                                          Length 21;
                                                                                                                                                                                                                                                          1; Indels
capture probes used in the method of the invention.
                                                                                Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                     Score 19.4; DB 1;
Pred. No. 7.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat DNA microarray capture oligonucleotide #48.
                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                      2708 TAAAAAAAAAAAAAAA 2728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                            21 TATAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 5; 8pp; German.
                                                                                                                                                                     0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK01328 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004 (first entry)
                                                                                                                                                        Query Match
Best Local Similarity 95.2°
Matches 20, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Boekenkamp D, Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-714082/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK01328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 71
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cagnits that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comparises anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises are particular sorts of single stranded nucleic acids. The capture capture of sequence of 10-50, particularly 15-25, T residues. The capture of comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, caranic, glass, cinteracting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (bysical, stimulated by an electrical filaments and binding of single compariation is used (i) for analysis of patterns, especially in mucosal, capture root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food
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      and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                        .
0
                                                                                                                                                                      0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; ive 0; Mismatches 1; Indels
                                                                                       capture probes used in the method of the invention
                                                                                                                                Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat DNA microarray capture oligonucleotide #57.
                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAAA 2729
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                                                                                                                                                                                                                                                                                                       21 AAGAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                 ADK01337 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                 Matches 20; Conservative
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                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rattus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK01337;
                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                        RESULT 715
                                                                                                                                                                                                                                                                                                                                                                                               ADK01337,
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additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. 8888888888888

Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

; 0 Score 19.4; DB 1; Length 21; Pred. No. 7.6e+02; 1; Indels 0; Mismatches 2708 TAAAAAAAAAAAAAAAAA 2728 0.7%; Conservative Query Match Best Local Similarity Matches 20; Conserv à

TGAAAAAAAAAAAAAAA 1 21 유

ADK01282 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01282; RESULT 716 

Rat DNA microarray capture oligonucleotide #2.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattue

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Dieck HT, Hoppe Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions. patterns and and and and

Example; Page 4; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a3 'variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular,

physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, activity of blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals analyor nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible, ADK01281-ADK01344 represent capture probes used in the method of the invention.

Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;

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Gaps

Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 55.2%; Pred. No. 7.6e+02; Indels 0; Mismatches 95.2%; 20; Conservative Similarity Query Match Best Local Matches

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ADK01334 standard; DNA; 21 RESULT 717 ADK01334/c 

BP.

ADK01334;

Rat DNA microarray capture oligonucleotide #54.

(first entry)

06-MAY-2004

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

2002DE-01008794. 28-FEB-2002; (DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Boekenkamp D, Dieck HT,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25. T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass,

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metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, chair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, organic additives or supplements, especially minerals, rance elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid bool. Since the method is based on comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the comparison of muchaic acid pools, not individual genes, matrix miniaturisation is possible. ADX01281-ADX01344 represent
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Sequence 21 BP; 0 A; 0 C; 2 G; 19 T; 0 U; 0 Other;

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0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; ive 0; Mismatches 1; Indels
                                                2707 CTAAAAAAAAAAAAAAAAA 2727
                                                                         21 CCAAAAAAAAAAAAAAAAA 1
                           Conservative
           Similarity
  Mar.
Local Sir.
 Query Match
                        Matches
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Gaps

ADK01296 standard; DNA; 21 BP (first entry) 06-MAY-2004 ADK01296; RESULT 718 ADK01296, 

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #16.

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Dieck HT, Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture

comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cp bysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the celivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible crandicular singing of the complete pattern of all nucleic acid in a cell, and can detect very small differences in pools of nucleic acid pool. Since the center probes used in the method of sols.

C method is based on comparison of nucleic acid pool. Since the center probes used in the method of the invention.

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Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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Gaps
                                ö
0.7%; Score 19.4; DB 1; Length 21; 55.2%; Pred. No. 7.6e+02; ve 0; Mismatches 1; Indels
             95.2%;
                             20; Conservative
               Similarity
 Query Match
                 Best Local
                              Matches
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ADK01338 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01338; ADK01338, 

Rat DNA microarray capture oligonucleotide #58.

capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. ss; hybridisation; blood; nerve; germ

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794.

Hoppe H; Dieck HT, Ď, Boekenkamp

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example, Page 6; 8pp, German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides

have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 indicatides to allow binding of particularly locked nucleic acids (LNA) and the anchor region agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the certivity of pharmaceuticals and/or nutritional compounds, esg food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acid pool; since the method is based on comparison of nucleic acid pool; since the method is based on comparison of nucleic acid pool; since the method is based on comparison of nucleic acid pool; since the capture probes used in the method of the invention. 8888888888888888888888888888888**8** 

Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; Indels ä 0; Mismatches Best Local Similarity 95.2 Matches 20; Conservative Query Match

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2707 CTAAAAAAAAAAAAAAAAA 2727

21 CGAAAAAAAAAAAAAAAA 1 ઠે g

ADK01320 standard; DNA; 21 BP ADK01320; RESULT 720 

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #40.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Dieck HT, Hoppe 3oekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable patterns and screening and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Hoppe H;

Boekenkamp D, Dieck HT,

WPI; 2003-714082/68.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794.

DE10208794-A1.

Rattus sp.

04-SEP-2003.

cc reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides converse as 1-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleic acids. The capture oligonuclectides for single stranded nucleic acids. The capture or comprises a sequence of 10-50, particularly 15-25, T residues. The capture of 10-50, particularly 15-25, T residues. The capture or capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single tranded nucleic acids to the surface is (quasi) conducting the surface is quasic, submalated by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, conditives or supplements, especially minerals, trace elements, organic acidis (anino, carboxyllor or farty acid) or thair derivatives. Grabox, nerve or germ cells and (ii) for determining the captures or supplements, especially minerals, trace elements, organic acidis and mixtures. The method provides rapid, inexpensive and reproducible conditions in an early and reproducible conditions in method is based on comparison of all nucleic acid pool. Since the capture and an acidi pool is the invarient or acidi pools of the complete pattern of all nucleic acid bools. Since the capture and an acidi pools of the invariant or the nucleic acid pool. Since the capture and aci ö ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Gaps ö Length 21; 1, Indels capture probes used in the method of the invention. Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other; Score 19.4; DB 1; Pred. No. 7.6e+02; Rat DNA microarray capture oligonucleotide #24. 0; Mismatches 2706 ACTAAAAAAAAAAAAAAA 2726 21 ACGAAAAAAAAAAAAAAA 1 BP. 0.7%; ADK01304 standard; DNA; 21 (first entry) 20; Conservative Query Match Best Local Similarity 06-MAY-2004 ADK01304; Matches RESULT 721 ADK01304/ g HARAK KARAK KARAN KARAK KARAK KARAK LA KARAK LA

XXX999999999999999999999999999999

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleocides compariable andhor regions. The capture oligonucleocides and non-variable regions. The capture oligonucleocides and solid matrix and (b) that comprises all possible combinations of up to 10 nucleocides to allow binding of particular sorts of single stranded nucleic acids. The capture of comprises a sequence of lock, particularly 15-25, T residues. The agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-60, particularly 15-25, T residues. The compuring properties are biotinylated and immobilised on a surface interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cation, carboxylic or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid inexpensive and reproducible caid and a call, and a call in a cell, and a call in a cell, and acids (amino, carboxylic or fatty acid) or their derivatives, salts and acids (amino, carboxylic or fatty acid) or their derivatives, salts and callows imaging of the complete pattern of allows in arrix miniarium is analysing a nuclei is possible. ADK01281-ADK01344 represent genes, matrix miniaturisation is possible. ADK01281 capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 0 C; 2 G; 19 T; 0 U; 0 Other;

.; 0 DB 1; Length 21; 1; Indels Query Match 0.7%; Score 19.4; DB 1; Best Local Similarity 95.2%; Pred. No. 7.6e+02; Matches 20; Conservative 0; Mismatches 1. 2706 ACTAAAAAAAAAAAAAAA 2726 21 ACCAAAAAAAAAAAAA 1 à 8

RESULT 722

ADK01325 standard; DNA; 21 BP

ADK01325;

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #45.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68 

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

Thus invention describes a novel method for sorting single-stranded controlled acids by isolation and hybridisation of nucleic acids pools, then reading out, where the nucleic acids are selectively bound using capture agence that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agence are particularly locked nucleic acids (LIAA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceramic, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, physical, sused (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cardis (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible candetect very small differences in the nucleic acids from cells, and can lower matrix, miniaturiar, in a call nucleic acid bools, since the complete pattern of all nucleic acid soil on the individual and can detect very small differences in the nucleic acid coid doil or eacid soil on the individual and can detect very small differ method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent invention describes a novel method for sorting single-stranded capture probes used in the method of the invention.

Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

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Gaps

Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; ve 0; Mismatches 1; Indels 95.2%; 20; Conservative Query Match Best Local Similarity Matches

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ઠે 셤 RESULT 723

ADK01292 standard; DNA; 21 BP ADK01292/

ADK01292;

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #12.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. ss; hybridisation;

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794. 

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H;

Dieck HT,

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uncleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides comprises and non-variable regions. The capture oligonuclectides comprises all possible combinations of up to 10 nucleotides to allow that comprises a sequence of single stranded nucleic acids. The capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceramic, resh, agel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface cation is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives.

Conducting properties in pools of nucleic acids from cell, and center the method is based on comparison of nucleic acid pools, not individual capture probes used in the method of the invention.
                                                                                                                              Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                     This invention describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                    Example; Page 5; 8pp; German.
                                                                                                                                                                                          and constant regions.
                                                                                WPI; 2003-714082/68
                         Boekenkamp D,
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0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels 2706 ACTAAAAAAAAAAAAAAA 2726 21 AGTAAAAAAAAAAAAAAA 1 Query Match
Best Local Similarity 95.2
Matches 20; Conservative

Rat DNA microarray capture oligonucleotide #32. ADK01312 standard; DNA; 21 BP 06-MAY-2004 (first entry) 

se; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

DE10208794-A1 Rattus sp.

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Boekenkamp D,

WPI; 2003-714082/68

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded cucles acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cadding out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to hat comprises all possible combinations of up to 10 nucleotides to allow binding of particular sortes of single stranded nucleic acids. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic glass, comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular flaments and binding of shingle cranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical fidal or through a molecular, physical, stimulated by an electrical fidal or through a molecular side activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, or for acids (anino, carboxylic or fatty acid) or through a molecular or activity of pharmaceuticals and/or nutritional compounds, e.g. food acids (anino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and repreducible cand sind can detect very small differences in the nucleic acids from cells, and can detect very small differences in the nucleic acid pool, and capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;

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Gaps

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Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; tive 0; Mismatches 1; Indels Best Local Similarity 95.2 Matches 20; Conservative Query Match

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RESULT 725

g 8

Rat DNA microarray capture oligonucleotide #18. ADK01298 standard; DNA; 21 BP. (first entry) 06-MAY-2004 ADK01298;

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. BXSXXXXXXXXXXXX

Rattus sp.

DE10208794-A1

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region, the complement of which is present at least once in each nucleic acid and a3 'variable, discriminatory region that comprises all possible combinations of up to 10 nucleocides to allow binding of particular sorts of single stranded nucleic acids. The capture or particularly locked nucleic acids (INA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture or particularly locked nucleic acids (INA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture or sequence of 10-50, particularly 15-25, T residues the capture conducting properties and especially in the form of a chip. Its surface interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, es, of food acids (amino, carboxylic or fatty acid) or though a plants of captures. The method provides rapid, inexpensive and reproducible representation of differences in pools of mucleic acid being acid in a cell, and enthod is based on compatison of nucleic acid pools, not individual genes, matrix miniaturisation is the method of the invention.
                                                                                                                                                                                                                                                                                                                            Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                              Hoppe H;
                                                                                                                                                                          (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                           28-FEB-2002; 2002DE-01008794.
                                                                                                                            28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 5; 8pp; German.
                                                                                                                                                                                                                         Boekenkamp D, Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                        and constant regions.
                                                                                                                                                                                                                                                                          WPI; 2003-714082/68
                         04-SEP-2003
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Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; 1; Indels Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other; 0; Mismatches Best Local Similarity 95.2 Matches 20; Conservative Query Match

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2707 CTAAAAAAAAAAAAAAAA 2727 21 CTCAAAAAAAAAAAAAA 1 ઠ g

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #56. ADK01336 standard; DNA; 21 BP. 06-MAY-2004 (first entry) ADK01336; RESULT 726 ADK01336/ 

(DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794 3oekenkamp D, Dieck HT, WPI; 2003-714082/68. DE10208794-A1 04-SEP-2003. Rattus sp. 

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture careading out, where the nucleic acids are selectively bound using capture careading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (in nucleotides to allow binding of particularly locked nucleic acids (in(A)) and the anothor region comprises a sequence of 10-50, particularity is-25. T residues. The agents are particularly locked nucleic acids (in(A)) and the anothor region comprises a sequence of 10-50, particularity is-25. T residues. The method is properties and especially in the form of a chip. Its surface is particularly a layer of (blo) molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo) molecular filaments and binding of single conducting propertical field or through a molecular, physical, stimulated by an electrical field or through a molecular, computed by an electrical field or through a molecular, content root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cailes (amino, carboxylic or fatty acid) or their derivatives, salts and cailows inaging of the complete pattern of all nucleic acid fool. Since the anlows inaging in the complete pattern of all nucleic acid pool. Since the cannot and an analysis of the complete pattern of acid pool. Since the cannot and in the mucleic acid pool. Since the capture and i Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable capture probes used in the method of the invention. Example; Page 6; 8pp; German. and constant regions.

Gaps .. 0 0.7%; Score 19.4; DB 1; Length 21; 95.2%; Pred. No. 7.6e+02; 1; Indels Sequence 21 BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other; 0; Mismatches 2706 ACTABABABABABABABABA 2726 20; Conservative Best Local Similarity Query Match Matches ઠે

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21 ACAAAAAAAAAAAAAA 1 ADW71579 standard; DNA; 21 RESULT 727 윱

21-APR-2005 (first entry) ADW71579; XXXXX

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The invention relates to a complex (CI) comprising a first and second single stranded nucleic acid or its derivative and a metal ion, where the first and second base of the strands forms a bond with metal ion. Also included are detecting the existence of thymine-thymine, cytosine.

Cytosine or cytosine-thymine base pair in a DNA or its analog (involving melting DNA or its analog in an aqueous medium, processing the solution of an acmoniations of Hg(II)2+ and Ag+ in the prepared DNA solutions, and comparing the characteristics of the solution, where change in characteristics of the solution, where change in characteristics of T.T. base pair, comparing the existence of T.T. base pair, comparing a metal ion (comprising one or more DNA molecules or their analogs having a metal binding region, where the coupling of metal ion is detected by analyzing the characteristic change in DNA). The complex (CI) is useful as a non-watson Crick base pair metal complex (CI) is useful as a non-watson Crick base pair in a double stranded DNA. The propales to detecting non-watson Crick base pair in a double stranded DNA. The propales to detect non-watson Crick base pair in a double stranded DNA. The propales of characteristic change in a double stranded DNA. The prepared DNA. The DN
                                                                                                                                                                                                                             /*tag= a
/note= "Paired with a thymine on the opposite strand via
Non-Watson-Crick base pairing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Complex useful for detecting non-Watson Crick base pair in double stranded DNA, comprises first and second single stranded nucleic acid or its derivative and metal ion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detect non-Watson Crick base pair in a double stranded DNA. The present sequence is a 21mer double stranded oligonucleotide with 1 Non-Watson-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 19.4; DB 1; Length 21; Pred. No. 7.6e+02; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 20 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 32; 73pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ААААААААТАААААААА 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUL-2004; 2004WO-JP010300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JUL-2003; 2003JP-00201500.
26-FEB-2004; 2004JP-00051320.
Oligonucleotide DS21mer(T-T).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AED42748 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-2005 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2005-162557/17.
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                                                           DNA detection; ds.
                                                                                                                                                                                                                                                                                                                                                 WO2005010177-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ONOA/) ONO A.
                                                                                                                    Unidentified
                                                                                                                                                                                                        misc feature
                                                                                                                                                                                                                                                                                                                                                                                                       03-FEB-2005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AED42748;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ono A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a complex (CI) comprising a first and second single stranded nucleic acid or its derivative and a metal ion, where the first and second base of the strands forms a bond with metal ion. Also included are detecting the existence of thymine-thymine, cytosine.

Cytosine or cytosine-thymine base pair in a DNA or its analog (involving melting DNA or its analog in an aqueous medium, processing the solution.

CC 3 minutes, to obtain three DNA solutions, dissolving Hg(II)2+, Ag+ and combinations of Hg(II)2+ and Ag+ in the prepared DNA solutions, and comparing the characteristics of the solution, where change in characteristics in Hg(II)2+, Ag+ and combinations of Hg(II)2+ and Ag+ represents the existence of T-T base pair in the respective DNA solutions) and an agent (AI) for detecting a metal ion (comprising one or more DNA molecules or their analogs having a metal binding region, where the coupling of metal ion is detected by analyzing the characteristic change in DNA). The complex (CI) is useful as anon-watson Crick base pair in a double stranded DNA. The complex (CI) enables to detect non-Watson Crick base pair in a double stranded DNA. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                      /*tag= a
/note= "Paired with a cytosine on the opposite strand via
Non-Watson-Crick base pairing"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Complex useful for detecting non-Watson Crick base pair in double stranded DNA, comprises first and second single stranded nucleic acid or its derivative and metal ion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence is a 21mer double stranded oligonucleotide with 1 Non-Watson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 20 A; 1 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUL-2003; 2003JP-00201500
26-FEB-2004; 2004JP-00051320
     Oligonucleotide DS21mer(C-C)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUL-2004; 2004WO-JP010300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-APR-2005 (first entry)
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Best Local Similarity
Marches 20; Conserv
                                                                 DNA detection; ds.
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                                                                                                                                                                                                           misc_feature
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Ono A;

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Gaps

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ADW71578;

X2X5X

RESULT 728

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WPI; 2005-664172/68.
         gene silencing.
                                          29-APR-2002;
                                         18-APR-2002
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Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive; Antidiabetic: Cardiant; Neuroprotective; Antidiabetic; Antiinflammatory; Antibacterial; Osteopathic; Anorectic; Virucide; Anti-HIV; Nephrotropic; Antiatteriosclerotic; Muscular-Gen.; protein interaction; protein microarray; cancer; familial adenomatous polyposis; protein microarray; cancer; familial adenomatous polyposis; gastrointestinal-gen.; ischemia; cerebrovascular ischemia; autoimmune disease; destropersis; obesity; viral infection; acquired immune deficiency syndrome; glomerulonephritis; atherosclerosis; muscular dystrophy; ss; short interfering RNA; RNA interference; Protein interacting gene transcript siRNA sense oligo #173.

Unidentified; Synthetic.

/note= "3' overhang comprising two 2'-deoxythymine residues linked by a 5'-3' phosphodiester linkage" Location/Qualifiers /note= "3' 10. .21 \*tag= misc\_feature

US2005222029-A1.

06-OCT-2005

07-MAR-2005; 2005US-00075234

2001US-0276179P. 2001US-0277013P. 2001US-0284095P. 2001US-028420P. 2001US-0284404P. 2001US-0287513P. 2001US-0304101P. 2001US-0307233P. 2001US-0347829P. 2001US-0259573P 16-APR-2001; 19-MAR-2001; .7-APR-2001; 30-APR-2001; 04-JAN-2001 9-APR-2001

2002US-0349843P. 2002US-0354899P. 2002US-00098979. 2002US-00099924. 2001US-0343818P. 2002US-00035344. 2002US-0346384P. 10-JUL-2001; 23-JUL-2001; 22-OCT-2001; 25-OCT-2001; 04-JAN-2002; 07-JAN-2002; 07-JAN-2002; 06-FEB-2002;

2002US-00100503 2002US-00124767 18-MAR-2002; 14-MAR-2002; LS-APR-2002; -APR-2002; 17-APR-2002;

(MYRI-) MYRIAD GENETICS INC.

2002US-00135802

Sugiyama J, Wettstein DA, Cimbora D, ď Bartel

Heichman K;

New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma

Disclosure, Fig 62, 198pp, English.

The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method for selecting modulators of the protein complex; a method of selecting modulators of an interaction between a first protein and a second protein; and the treating and/or preventing of diseases and disorders associated

ö assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders associated with the protein complexes. The diseases and disorders include cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, osteoporosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an siRNA oligo which targets the transcript of a protein forming part of a protein-protein complex of the invention. with the protein complexes. The protein complexes are useful in screening Gaps ö 0.7%; Score 19.4; DB 1; Length 21; 76.2%; Pred. No. 7.6e+02; tive 4; Mismatches 1; Indels Sequence 21 BP; 5 A; 7 C; 3 G; 2 T; 4 U; 0 Other; 1063 CCACGGATCTGACTACTCACT 1083 CCACGGAUCUGACUACUCATT 21 16; Conservative Local Similarity Query Match Matches ò 셤

DNA probe used in fingerprinting technique. BP AAT68615 standard; DNA; 24 (first entry) 20-FEB-1998 AAT68615; RESULT 730 AAT68615/c 

probe; screening; fingerprinting; assay; 3' termini; hybridisation; ss. 96EP-00118921 26-NOV-1996; EP778351-A2 11-JUN-1997. Synthetic.

Vematsu C; Kambara H, Okano K, (HITA ) HITACHI LTD WPI; 1997-300347/28

95JP-00311949

30-NOV-1995;

Nucleic acid assay methods - based on restriction fragment length determination.

Example 1; Page 7; 21pp; English.

The present sequence is a DNA probe used in a novel method of analysis or assay for nucleotides, which comprises: (i) digesting DNA with a castriction enzyme; (ii) discriminating a difference in sequences of the DNA fragments obtained around the 3' termini with a DNA probe and extending the DNA probe by a complementary strand synthesis to fractionate the DNA fragments into groups, and (iii) measuring lengths of the DNA fragments which belong to the groups, or length of the extended DNA probe, and using the lengths obtained for the fragments around the 3' termini as fingerprints. Where polyA is present, the presence of recognition sequence GCG is critical for clarifying the terminal site, this is because the length of polyA cannot be controlled. The method is useful for assaying long DNA sequences

Sequence 24 BP; 0 A; 2 C; 1 G; 19 T; 0 U; 2 Other;

Score 19.4; DB 1; Length 24; Pred. No. 8.1e+02; 0.7%; Best Local Similarity Query Match

Tue Nov

Matches

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The present invention relates to the isolation of polypeptide-laminin B210.67, and the polymucleotide encoding it. Also described is the process for preparing the protein by DNA recombination. The polypeptide is useful for treating diseases such as embryo development tratogenesis. The present sequence for reverse transcriptuse (RT)-PCR primer #1 is used with RT-PCR primer #2 (ABK12410) for isolating cDNA encoding polypeptide-laminin B210.67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polypeptide-laminin B210.67, useful for treating diseases such as embryo
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"this base represents an unspecified number of
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"this base represents an unspecified number
                                                                                       RT-PCR primer #1 for cDNA encoding polypeptide-laminin B210.67.
                                                                                                                           Polypeptide-laminin B210.67; embryo development teratogenesis; cytostatic; reverse transcriptase-PCR; RT-PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Replication error; drug development; somatic instability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fragment of a plasmid used to detect somatic instability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 BP; 19 A; 2 C; 0 G; 3 T; 0 U; 0 Other;
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0.7%; Score 19.4; DB 1;
Best Local Similarity 95.2%; Pred. No. 8.1e+02;
Matches 20; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                 (BODE-) BODE GENE DEV CO LTD SHANGHAI.
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                                                     (first entry)
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bases"
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misc_feature
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                                                                                                                                                                                         Unidentified
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               ABK12409;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polymucleotide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allelectused in a hybridisation assay for determining the identity of a nucleotide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Early-stage diagnosis of prostate cancer relies on prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its inable to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure which would enable early-stage diagnosis, and for preventative and curature sequences of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease
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ö
                                                                                                                                                                                                                                                                                                                           PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of a prostate cancer associated gene and biallelic markers derived
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Gaps
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 Indels
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Mismatches
                                                                                                                                                                                                                                                                                         PCR primer PGRT32 for PG1 coding sequence.
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98US-0099658P.
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                                                                                                                                                                                                                                                     (first entry)
   Conservative
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hes 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                     PSA; human; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-DEC-1998;
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09-SEP-1998;
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Length 24;

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Query Match

Best Loca Matches

ઠે 요 RESULT 732

ABK12409 ID ABK1

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The present invention relates to a method for determining if a gene product or compound is involved in preventing replication error in a cell. The method involves providing a cell with a specific inhibitor for a gene product or with a compound and determining the expression level of a marker gene in the cell, where the expression level of the marker gene is dependent on the occurrence of a replication error. The invention is useful in gene therapy and for treating a subject having tumours or cancer. The present sequence is a Caenorhabditis elegans heat-shock promoter DNA. This sequence is used to illustrate the method of
                                                                                                                                                                                                                                      Determining if a gene product/compound is involved in preventing replication error in a cell, useful for treating cancer, comprises determining expression level of a marker gene in a cell treated with a gene product inhibitor/compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 19.4; DB 1; Length 24; 87.0%; Pred. No. 8.1e+02; artive 0; Mismatches 3; Indels
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/note= "OTHER= locked nucleic acid"
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                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 3; 25pp; English.
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                22-MAY-2001; 2001EP-00201936.
22-MAY-2002; 2002WO-NL000322.
28-NOV-2002; 2002WO-WO095071.
                                                                                                                                                         Tijsterman M, Plasterk RHA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC48482 standard; DNA; 21
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nes 20; Conservative
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PLASTERK R H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention.
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                                                                                               (TIJS/)
(PLAS/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                               The specification describes a method for determining whether a product of a gene is involved in preventing a replication error in a cell. The method comprises providing the cell with a specific inhibitor for the product and determining the level of functional expression of a marker gene in the cell, where the level of expression of the marker gene is dependent on the occurrence of the replication error. The method is used for determining whether a product of a gene is involved in preventing a replication error in a cell. The identified genes are useful for developing diagnostic tools, or as targets for drug development to manipulate cells on the basis of the presence or absence of function of the gene. ABZ2335-36 represents fragments of plasmids used to detect somatic instability, in the course of the invention
                                                                                                                                                                                                                                                                                                             Determining whether a product of a gene is involved in preventing a replication error in a cell comprises providing a specific inhibitor for the product and determining the level of expression of a marker gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Best Local Similarity 87.0%; Pred. No. 8.1e+02;
Matches 20; Conservative 0; Mismatches 3; Indels
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"N can be repeated Y times"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Caenorhabditis elegans heat-shock promoter DNA #2.
                                                                                                                                                                        (NEVW-) KONINK NEDERLANDSE AKAD VAN WETENSCHAPPE
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                                                                                                                                                                                                                                Plasterk RHA, Tijsterman M;
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/note=
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                                                                                                                                                                                           (TIJS/) TIJSTERMAN M.
                                                                                                                                                                                                                                                                          WPI; 2003-129440/12.
            WO200295071-A2
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Query Match

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ADR44221;

RESULT 734 ADR44221

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Gaps

Sequence 21 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 2 Other;

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The present sequence is that of pyrene-anchored locked nucleic acid (LNA) coligo(dT) primer ON12, which was used in first-strand cDNA synthesis from enkaryotic mRNA. It includes compound '17d' at its 3' end, which is based on an LNA-type 2'-0,4'-C-methylene-beta-D-ribofuranosyl moiety. It is one of a set of such primers (see also ACC48483-85) that were used in an example from the invention to demonstrate improved reverse transcription of mRNA using pyrene-LNA anchored oligo(T) primers. The following results were observed: efficient priming on mRNAs with short poly(A) tails; efficient anchoring of the oligo(T) primer by pyrene-LNA and LNA-C/G/T units resulting in an improved T20-VN anchor primer and thus avoiding reverse transcription of long poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; and improved reverse that comprise unique base groups. Desirable nucleobase and nucleosidic that comprise unique base groups. Desirable nucleobase and nucleosidic into nucleic acid strands. The novel LNA compounds can be used e.g. as come in diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel nucleic acid comprising a locked nucleic acid unit having a modified base that comprises an optionally substituted carbocyclic aryl moiety, or modified nucleobase or nucleosidic base other than
                   "mod_base= OTHER
'note= "OTHER= locked nucleic acid"
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/mod_base= OTHER
/note= "OTHER= locked nucleic acid"
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/mod_base= OTHER
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22-SEP-2001; 2001US-0323967P.
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Determining the relative copy number of a group of target nucleic acid molecules present in a sample by performing a first or second PCR in a PCR mixture and quantifying the number of copies of the second target nucleic acid product.
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                                   Gaps
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                                                                                                                                                                                                                                                                                                             Multiplex real-time quantitative PCR; PCR primer; copy number; Alzheimer's disease; 85.
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 Length 21;
                                 Indels
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0.7%; Score 19.2; DB 1;
95.0%; Pred. No. 7.9e+02;
tive 1; Mismatches 0;
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19-JUL-2002; 2002US-0397475P.
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                                                                                                                                                                                   ACC99729 standard; DNA; 21
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                                      Conservative
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                      Best Local Similarity
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                                                                                                                                                                                                                                                                                  Oligonucleotide
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                                      19;
                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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     Query Match
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Matches
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Homo sapiens
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                                 Human;
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                                                                                                                     Vaccine, cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or stabhylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a the control of the present and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                       Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
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llarity 87.5%; Pred. No. 8.4e+02;
Conservative 0; Mismatches 3
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                                                                                                 Immunostimulatory nucleic acid #51.
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                              AAF98935 standard; DNA; 24 BP.
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23-AUG-2000; 2000US-0227436P.
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(COLE-) COLEY PHARM GMBH.
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                                                                          (first entry)
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                                                                          12-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21;
                                                                                                                                                                                Synthetic.
                                                   AAF98935;
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Best Local
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                    AAF98935,
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Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; labroplasia; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated polypeptide of human tre carcinogenic gene protein 10.56 comprising a 96 residue amino acid sequence, fully defined in the specification, or its fragment, analogue or derivative. The polypeptide is useful in the diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) present sequence is a primer used to amplify a polynucleotide encoding the polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human tre carcinogenic gene protein 10.56 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
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                                                                           Tre carcinogenic gene protein 10.56; cytostatic; haemostatic;
                                                                                                                 virucide; immunomodulatory; antiinflammatory; gene therapy; cancer; haemopathy; human immunodeficiency virus; HIV; infection; immunological disease; inflammatory disorder; PCR primer; ss.
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37.5%; Pred. No. 8 48+02;
Ve 0; Mismatches 3; Indels
Human Tre carcinogenic gene protein 10.56 PCR primer 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAY-2000; 2000CN-00115824.
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21; Conservative
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WPI; 2002-075668/10.

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                                                                                                                                                                                                                                                                                                                  administering at least one antitanglogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antianglogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumaroid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myccardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                     Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                       comprising
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                                                                                                                                                                                                                                                                                                      invention relates to inhibiting angiogenesis in a subject,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Score 19.2; DB 1; Length 24;
87.5%; Pred. No. 8.4e+02;
ative 0; Mismatches 3; Indels
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                                                                                                                    (COLE-) COLEY PHARM GROUP INC.
                                                                                  2000US-0255534P.
                                               14-DEC-2001; 2001WO-US048458.
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nes 21; Conservative
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                                                                                 14-DEC-2000;
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                                                                                                                                                     Bratzler RL;
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                 11-JUL-2002
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                                                                                                                                          This invention describes a novel human tre oncogene 10.56 which has cytostatic, haemostatic, virucide, immunomodulatory and antiinflammatory activity and can be used for gene therapy. The polypeptide of the invention and its encoding polynucleotide are used in diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. Sequence represents an RT-PCR primer used in the amplification of the human tre oncogene 10.56 gene which is described in the disclosure of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human transcriptional activation subunit 14 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                            human tre oncogene 10.56 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; transcriptional activation subunit 14; malignant neoplasm; haematopathy; cytostatic; HIV infection; human immunodeficiency virus; immunological disease; inflammation; virucide; immunomodulatory; antiinflammatory; reverse transcriptase-PCR; RT-PCR; primer; ss.
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0.7%; Score 19.2; DB 1;
Best Local Similarity 87.5%; Pred. No. 8.46+02;
Matches 21; Conservative 0; Mismatches 3;
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                                                                                                            Example 2; Page 12; 32pp; Chinese
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                                                                                                                                                                                                                                                                                                          invention
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BP.

(first entry)

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Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                 ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                             Immunostimulatory nucleic acid #51.
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                                                                                            ADB36437 standard; DNA; 24
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(PETE/) PETERSEN D
(FOUR/) FOURON Y.
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                                     RESULT 743
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various inflammations. Antagonists against the polypeptide can also be used in treating such diseases. The present sequence for reverse transcriptase (RT)-PCR primer #2 is used with RT-PCR primer #1 (ABK13714) for isolating cDNA encoding human transcriptional activation subunit 14
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                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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                                                                                                                                                                                        Query Match 0.7%; Score 19.2; DB 1; Length 24; Best Local Similarity 87.5%; Pred. No. 8.4e+02; Matches 21; Conservative 0; Mismatches 3; Indels
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                                                                                                                                       Sequence 24 BP; 0 A; 2 C; 2 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACD99368 standard; DNA; 24
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The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.
                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                Match 0.7%; Score 19.2; DB 1; Length 24; Local Similarity 87.5%; Pred. No. 8.4e+02; es 21; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                               Sequence 24 BP; 0 A; 0 C; 3 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAAAAAAA 2732
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24 AAAAACAAAAAAACAAAAAAAA 1

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                                                                                                                                                                          New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
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87.5%; Pred. No. 8.4e+02;
tive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                         Claim 14; SEQ ID NO 27; 139pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG75926 standard; DNA; 24 BP
                                                   30-MAY-2003; 2003WO-EP005691.
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                                                                          30-MAY-2002; 2002CA-02388049
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                                                                                                   (IMMU-) IMMUNOTECH SA
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Matches 21; Conserv
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WO2003101375-A2
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                                                                                                                                                                                                                    melanoma
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This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA cligonucleotides (without a CpG motif), which can stimulate an immune cresponse in animals of the order of primate, including humans. The immune cresponse is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present cinvention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell continum and this oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo incare asserved.
                                                                                                                                                                                                   New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
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Pred. No. 8.4e+02;
0; Mismatches 3;
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Local Similarity 87.5%;
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                                                                                                                                                                                                                                                                                                                             melanoma
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New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. Claim 14; SEQ ID NO 24; 139pp; English melanoma

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CPG motif.) which can stimulate an immunne response in animals of the order of primate, including humans. The immune and antibody production in Be-ealls, as well as cell differentiation, cytokine cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an entigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or cells and sylocolipids. As used in activation, treating, preventing or therapy for inducing B-cell activation, treating preventing or calls and elogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo, used in an exemplification of the invention.

Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 19.2; DB 1; Length 24; 87.5%; Pred. No. 8.4e+02; ive 0; Mismatches 3; Indels 24 AAAAAAAACAAAATGAAAAAA 1 Query Match
Best Local Similarity 87.5'
Matches 21, Conservative

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Immunostimulatory non-CpG oligonucleotide IMT 179 SegID 26. ADG75924 standard; DNA; 24 BP. 11-MAR-2004 (first entry) ADG75924;

se; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differantiation; production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;

renal cell carcinoma

WO2003101375-A2.

11-DEC-2003

30-MAY-2003; 2003WO-EP005691.

30-MAY-2002; 2002CA-02388049.

(IMMU-) IMMUNOTECH SA.

Lopez RA;

WPI; 2004-053333/05.

New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. melanoma

Claim 14; SEQ ID NO 26; 139pp; English

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CPG motif.) which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, virtues, bacteria, parasites, tumour cells and glycoligids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi, sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo, used in an exemplification of the invention.

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Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;

Gaps .. 0 0.7%; Score 19.2; DB 1; Length 24; 3; Indels Pred. No. 8.4e+02; 0; Mismatches 3 87.5%; Ouery Match
Best Local Similarity 87.57
Warches 21; Conservative

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24 AAAAACAAAATGAAAAAAAAAAA

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Non-CpG DNA oligonucleotide 2. ADG76001 standard; DNA; 24 BP (first entry) 11-MAR-2004 ADG76001; RESULT 748 ADG76001

ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma renal cell carcinoma. 

Synthetic.

WO2003101375-A2.

11-DEC-2003.

30-MAY-2003; 2003WO-EP005691.

30-MAY-2002; 2002CA-02388049.

SA. (IMMU-) IMMUNOTECH

Lopez RA;

WPI; 2004-053333/05.

New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. melanoma.

Example 17; Page 80; 139pp; English.

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindronic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present

Sequence 24 BP; 0 A; 0 C; 3 G; 21 T; 0 U; 0 Other;

invention

ö This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune and antibody production in B-cells, as well as cell differentiation, cytokine and antibody production in Daemacycoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or therapy for inducing B-cell activation, treating, preventing or ameniocating an immune system disporder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is a non-CpG DNA oligo of the antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system discorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is a non-CpG DNA oligo of the New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g. ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; invention describes immunomodulator compositions that also comprise an Gaps ö 0.7%; Score 19.2; DB 1; Length 24; ilarity 87.5%; Pred. No. 8.4e+02; Conservative 0; Mismatches 3; Indels Seguence 24 BP; 0 A; 0 C; 3 G; 21 T; 0 U; 0 Other; 2709 AAAAAAAAAAAAAAAAAAA 2732 Example 17; Page 81; 139pp; English. AAAAACAAAAAACAAAAAAAAA 1 Non-CpG DNA oligonucleotide 36. 30-MAY-2003; 2003WO-EP005691. 30-MAY-2002; 2002CA-02388049 ADG76035 standard; DNA; 24 (first entry) (IMMU-) IMMUNOTECH SA renal cell carcinoma. WPI; 2004-053333/05. Local Similarity nes 21, Conserv WO2003101375-A2 11-MAR-2004 11-DEC-2003 invention. Synthetic. 24 ADG76035; melanoma. Query Match Best Loca Matches ADG76035/ 8888888888888 ઠે 셤 

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Length 24;
                                                                                                                                                                                                                                                                                            Immunostimulatory non-CpG oligonucleotide IMT 174 SeqID 21.
                                   3; Indels
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 DB 1;
 Score 19.2; DB 1;
Pred. No. 8.4e+02;
                                   Mismatches
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Best Local Similarity 87.5%;
Matches 21; Conservative
                                                                                                                                                                                           24
                                                                                                                                                                                                                                                               (first entry)
                                     Conservative
                                                                                                                                                                                           ADG75919 standard; DNA;
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                   Local Similarity
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   Query Match
                                                                                                                                                           RESULT 750
                                     Matches
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This invention relates to novel immunostimulatory oligonuclectides that contain a non-palindromic sequence motif. Specifically, it refers to DNA coligonuclectides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine production in B-cells, as well as cell differentiation and antibody production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory phosphorothioate non-cpG variant DNA oligo, used to determine the effect of converse on a converse on an immunostimulatory converse on a converse on an immunostimulatory converse on a conve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunostimulatory non-CpG phosphorothioate DNA oligo IMT179 SeqID73.
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2709 AAAAAAAAAAAAAAAAAA 2732
                                                           exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG75971 standard; DNA; 24 BP
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New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.

Claim 14; SEQ ID NO 22; 139pp; English.

melanoma

ss; non-CpG; immunostimulatory; non-palindromic; immune response; ptollferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;

renal cell carcinoma

WO2003101375-A2

Synthetic

11-DEC-2003.

30-MAY-2003; 2003WO-EP005691. 30-MAY-2002; 2002CA-02388049.

SA.

(IMMU-) IMMUNOTECH

Lopez RA;

WPI; 2004-053333/05.

Immunostimulatory non-CpG oligonucleotide IMT 175 SeqID 22.

(first entry)

11-MAR-2004

ADG75920;

ADG75920 standard; DNA; 24 BP.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
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Matches
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Gaps

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0.7%; Score 19.2; DB 1; Length 24; ilarity 87.5%; Pred. No. 8.4e+02; Conservative 0; Mismatches 3; Indels

Query Match Best Local Similarity Matches 21; Conserv

2709 AAAAAAAAAAAAAAAAAAAA 2732

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24 AAAAACAAATGAAAAAAAA 1

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New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.
                                                                                                                                                                                                                                                                                                                                                                    Claim 14; SEQ ID NO 23; 139pp; English.
                                                                                                                                         30-MAY-2003; 2003WO-EP005691.
                                                                                                                                                                       30-MAY-2002; 2002CA-02388049.
                                                                                                                                                                                                                                                               WPI; 2004-053333/05.
                                                                                                                                                                                                   (IMMU-) IMMUNOTECH
                                                                               WO2003101375-A2.
                                                                                                             11-DEC-2003,
                                                  Synthetic
                                                                                                                                                                                                                               Lopez RA;
                                                                                                                                                                                                                                                                                                                                          melanoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune artibody production in B-cells, as well as cell differentiation, cytokine antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an invention describes immunomodulator compositions that also comprise an categor selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or mellorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunatory non-CpG
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                                                                                                                                                                                                                                                                                                                                                                                                 New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      se; non-CpG; immunostimulatory; non-palindromic; immune response;
proliferation; differentiation; cytokine; antibody production; B-cell;
plasmacytoid dendritic cell; immunomodulator; gene therapy;
                                                                ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        variant DNA oligo, used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.7%; Score 19.2; DB 1; Length 24;
87.5%; Pred. No. 8.4e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulatory non-CpG oligonucleotide IMT 176 SeqID 23.
                                  Immunostimulatory non-CpG oligonucleotide IMT 178 SeqID 25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 14; SEQ ID NO 25; 139pp; English.
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          11-MAR-2004 (first entry)
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hes 21; Conservative
                                                                                                                                                                                                                                                                                                               SA
                                                                                                                                renal cell carcinoma
                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-053333/05
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                                                                                                                                                                                        WO2003101375-A2.
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                                                                                                                                                            Synthetic.
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Matches
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This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif.) which can stimulate an immunne response in animals of the order of primate, including humans. The immune cresponse is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present cinvention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ampliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell correction and an immunostimulatory non-CpG variant DNA olice assets.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene typing; polymorphic microsatellite loci; PML; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             variant DNA oligo, used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%; Score 19.2; DB 1; Length 24; 87.5%; Pred. No. 8.4e+02; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cow prion protein microsatellite locus primer #88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
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                                                                                                                                                 Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                 Example 3; Page 29; 64pp; German.
                                                                                                  Han Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             microsatellite; PCR; primer; ss
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                          09-AUG-2002; 2002DE-01036711.
                                               09-AUG-2002; 2002DE-01036711.
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Best Local Similarity 87.5'
Matches 21; Conservative
                                                                                                  Geldermann H, Preuss S,
                                                                         (UYHO-) UNIV HOHENHEIM
                                                                                                                         WPI; 2004-215730/21.
                                                                                                                                                             identifying predispolength of amplicons.
26-FEB-2004.
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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA sample containing at least one segment of (1); and cast eremining the length of the resulting amplicon(8). Also described are: The method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or compare PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a diseases and for prediagnosis of such diseases, especially prion diseases to diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, cometabolic diseases; also to type genes that encode milk proteins, cometabolic diseases, also to type genes that encode milk proteins, particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion cycle protein (PrP) comprising a polymorphic microsatellite locus.
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                                                                                                                                                                                                                                           Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.7%; Score 19.2; DB 1; Length 24; Best Local Similarity 87.5%; Pred. No. 8.4e+02; Matches 21; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 0 A; 3 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Allergic response suppressor oligonucleotide #60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAAAAAA 2732
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bacterial infection; viral infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24 AAAAAAAAAGAAAGAAAGAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; Page 28; 64pp; German.
                                                                                 Han Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADU89376 standard; DNA; 24 BP
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                                                                                 Preuss S,
(UYHO-) UNIV HOHENHEIM.
                                                                                                                                                              WPI; 2004-215730/21.
                                                                                                                                                                                                                                                                                                                             length of amplicons.
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(PETE/) PETERSEN D
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2004235774-A1.
                                                                             Geldermann H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-FEB-2005
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Gaps

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WPI; 2004-833006/82

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The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the
                             Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; mimune stimulation; inflammation; psoriasis; inflammatory bowel disease; crohns disease; ulcerative colitis; eczema; skin allergy; contact dermatitis; ss; phosphorothioate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                  Sequence 24 BP; 0 A; 0 C; 3 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory oligonucleotide, SEQ ID 54.
                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                      Disclosure; SEQ ID NO 60; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAAAAAAAAAAA 2732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24 AAAAACAAAAAAAAAAAAAAAA 1
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87.5%;
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29-MAR-2002; 2002US-00112653
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                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.77
Best Local Similarity 87.5
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                    method of the invention.
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2005250726-A1
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modified_base
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The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating nonallergic inflammatory diseases such as psoriasis, inflammatory bowel diseases and ulcertive colitis), eczema, allergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                Score 19.2; DB 1; Length 24; Pred. No. 8.4e+02;
                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 19 BP; 1 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                            Sequence 24 BP; 0 A; 0 C; 3 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
              Disclosure; SEQ ID NO 54; 58pp; English
                                                                                                                                                                                            contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                         24 AAAAACAAAAAACAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВЪ
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                                                                                                                                                                                                                                                                Query Match

Best Local Similarity 87.5%;
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ75551 standard; DNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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Gaps

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0.7%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 7.7e+02; iive 0; Mismatches 0; Indels

Query Match
Best Local Similarity 100.
Matches 19; Conservative

2707 CTAAAAAAAAAAAAAA 2725

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The invention relates to aminooxy-modified(oligo)nucleotides or nucleosides which are useful as therapeutics, diagnostics, and research reagents. They may be used, e.g., for modulation of the ras gene and may be able to modulate the process of transformation from normal to malignant cell growth. They may be prepared using known methods. Inclusion of the aminooxy moleties can improve binding of oligonucleotides to complementary strands. The moieties can also provide conjugation sites useful for conjugation of useful ligands (e.g. reporter groups and groups for modifying uptake, distribution or other pharmacodynamic properties) to oligonucleotides. The present sequence in prepagents an example of an aminooxy-modified oligonucleotide disclosed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New aminooxy-modified oligonucleotides - which can show improved binding to complementary strands and improved resistance to nuclease.
                                                                                 phosphorothioate; ras gene; malignant cell growth; aminooxy-modified;
nuclease resistance; reporter group; ss.
                                                                                                                                                                                                                          /*tag= a
/note= "5-methyl, 2'-aminooxyethoxy-thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide containing modified internucleotide linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 7.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 84; 131pp; English.
                                                                                                                                                                             Location/Qualifiers
15. .18
                                          Aminooxy-modified oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAAA 2727
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98US-00016520
      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cook PD, Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-568232/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide; ss.
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modified_base
                                                                                                                                                                                                                                                                                 WO9835978-A1
        14-DEC-1998
                                                                                                                                                                                                                                                                                                                                                             13-FEB-1998;
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                                                                                                                                                                                                                                                                                                                        20-AUG-1998
                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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AAV06820/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
      셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequences given in AAT10742-67 are synthetic oligonucleotides which device (ED) of the invention of the electronically self-addressable device (ED) of the invention. The ED comprises a substrate, an electrode or opt. a number of electrodes supported by the substrate, a current square operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding to the electrode. The attachment layer is capable of attaching a macromolecule. The ED is used for genetic typing and comprises a number of electronically addressable locations each comprising an electrode, and a binding entity, such as one of these probes, attached to each of the locations capable of a genetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Electronically self-addressable device - used for electronic control of, e.g. nucleic acid hybridisation.
                                                                                                                                                                                                                                                     Electronically self-addressable device, ED; electrode; current source; attachment layer; permeable; counterion; genetic typing; probe; detection; ss.
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Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sosnowski RG
                                                                                                                                                                                                                                                                                                                                                                                                                                         -amino terminus"
                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                 757/c
AAT10757 standard; RNA; 19 BP.
CTAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tu E, Evans GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-00271882
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ID AAV07878 standard; DNA; 19
                                                                                                                                                                                                              Oligonucleotide probe, T-2
                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/note= "5'.
                                                                                                                                                                           09-SEP-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-1995;
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25-JAN-1996

Synthetic

AAT10757;

RESULT 760

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AAT10757,

Heller MJ,

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Gaps ö

Location/Qualifiers

Key

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RESULT 761

AAV07878;

SXXX

97US-00986065

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05-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss.
                                                                                                                                                                                                                                                            The invention relates, inter alia, to a method of preparing an oligonuclectide by coupling (1) a new nucleoside having a protected 5'-hydroxy group and at the 3'-position a group of formula -CH2-P(OR3)-NR4RS, with (2) a nucleoside or oligonucleotide having a free 5'-hydroxy group, to give (3) a precursor having an internucleoside linkage of formula -CH2-P(OR3)-O-; and converting this to a linkage of formula -CH2-P(OR3) (=X)-O- (where X = S or O). The present sequence is a specific
                 /note= "these T residues are formed as part of a conventional phosphoramidite oligonuclectide synthesis process but using as the reactant a thymosine nucleoside having at the 3'-position a group of formula -CH2-P(OCH2CH2CN)-N(iPF)2"
                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                          .
0
                                                                                                                                                                                                               New tetra:hydro:furan derivatives - useful in the synthesis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/note= "amino group attached at 5' terminal"
                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                           Douglas ME;
                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                  example of an oligonucleotide so prepared
                                                                                                                                                                           Altmann K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5' amino oligonucleotide probe T-2,
                                                                                                                                                                                                                                            Example 12; Page 29; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                                               19 AAAAAAAAAAAAAA 1
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                                                                                                                   97WO-GB001490
                                                                                                                                      96GB-00012600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX81316 standard; DNA; 19
                                                                                                                                                                           Collingwood SP, Moser HE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-AUG-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 19, Conservative
             /*tag= a
                                                                                                                                                                                              WPI; 1998-052233/05.
                                                                                                                                                         (NOVS ) NOVARTIS AG
                                                                                                                                                                                                                           oligo:nucleotide(s)
  modified_base
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                                                                              WO9747636-A2
                                                                                                                   03-JUN-1997;
                                                                                                                                      13-JUN-1996;
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                                                                                                18-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX81316;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX81316,
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The specification describes a self-addressable, self-assembling microelectronic device which is designed to actively carry out and control multi-step and multiplex molecular biological reactions in microscopic formate. A key aspect of this inventions is played by the ion microscopic formate. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation layer which overlies the electrode. This permeation control reaction but also separates the attached oligonucleocides and hybridized tometic mediately at the electrode surface. The microelectronic device is immediately at the electrode surface. The microelectronic device is mediately at the electrode surface. The microelectronic device is as nucleic acid hybridizations, antibody/antigen reactions, sample preparation, diagnostics and biopolymer synthesis. The device can electronically control the transport and attachment of specific microcof locations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific micro-locations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific micro-locations of evice is able to concentrate analytes and reactants, remove nonceptically bound molecules, provide stringency control for DNA bybridization reactions and improve the detection of analytes. The present sequence represents a probe used to exemplify the invention
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quenched fluorescent moiety; biological assay; detection; identification;
microorganism; sterilization assurance; nuclease; ss.
                                                                                                                                                                                   multi-step
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     a
"amine-modified C6 derivative of deoxythymidine
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"amine-modified C6 derivative of deoxythymidine
                                                      Edman CF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                      New microelectronic device quesigned to the sections in microscopic format and multiplex molecular biological reactions in microscopic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
                                                                                                                                                                                microelectronic device designed to carry out and control
                                                         Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
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100.0%; Pred. No. 7.7e+02;
1ve 0; Mismatches 0;
                                                         Nerenberg MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polynucleotide strand with amino groups.
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                                                                                                                                                                                                                                                                             Example 1; Page 90; 179pp; English.
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                                                            Tu E,
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                                                            WF
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/note=
(dT)"
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/note=
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(NANO-) NANOGEN INC.
                                                                                                                         WPI; 1999-385567/32.
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                                                                Sosnowski RG,
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AAX81927/
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The invention relates to a mammalian PG1 gene and protein, and a set of PG1 biallelic markers. The PG1 polynuclectide and biallelic markers are used in a hybridisation assay, a sequencing assay, or in an allelespecific amplification assay for determining the identity of a nuclectide at a PG1-related biallelic marker. The methods can be used to detect and to assess the risk of developing cancer or prostate cancer. Early-stage diagnosis of prostate cancer relies on prostate specific antigen (PSA) dosage. However, the effectiveness of this is limited due to its inability to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure which would enable early-stage diagnosis and for preventative and curative treatments of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligomeric compound; 2'-O-modified ribosyl nucleoside; 3' endo geometry; nuclease resistance; phosphodiester; ss.
                                                                                                                                                                                                                                                  Use of a prostate cancer associated gene and biallelic markers derived from it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                                                                                                                                                        Bougueleret L;
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-ive 0; Mismatches
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                                                                                                                                                                     Chumakov I,
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'note= "2'-modified
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/note= "2'-modified
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/note= "2'-modified
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                                                                                                                                                                                                                                                                                                                Claim 4; Page 374; 385pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAAA 2727
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                           98WO-IB002133
                                                               97US-00996306
98US-0099658P
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Best Local Similarity 100.(
Matches 19; Conservative
                                                                                                                                                                   Cohen D, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
/note=
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                                                                                                                             (GEST ) GENSET
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modified_base
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                           22-DEC-1998;
                                                                 22-DEC-1997;
                                                                                       09-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The specification describes an enzyme-specific cleavable polynucleotide substrate bearing quenched fluorescent moieties. The enzyme-specific cleavable polynucleotide substrate is useful in biological assays for detection and identification of microorganisms, sterilization assurance, pharmaceutical discovery, enzyme assays, immunoassays and other biological assays. The method provides a rapid and convenient approach for detection and identification of microorganisms. It can be adapted to sequence-dependent or sequence-independent tests. The invention provides improved accuracy, faster detection, and overall lower cost in detection and identification of microorganisms. The presence of nuclease is measured more accurately and sensitively by red-shifting the emission the electromagnetic spectrum and reducing the effect of background signal levels of intact reagents. The present sequence is used in the course of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen; PSA; human; ss.
'note= "amine-modified C6 derivative of deoxythymidine (dT)"
                                                           d
"amine-modified C6 derivative of deoxythymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                   enzyme-specific cleavable polynucleotide substrate bearing quenched
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer for PG1 biallelic marker 4-4-187.
                                                                                                                                                                                                                                                                                                       (MINN ) MINNESOTA MINING & MFG CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 20; 34pp; English.
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                                                           /*tag=
/note= "
(dT)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                        fluorescent moieties.
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                                                                                                                                                                                                                                                                                                                                                  Wei A, Mach PA;
                                          modified base
                                                                                                                                                                                                                         20-AUG-1998;
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                                                                                                                                            WO9935288-A1
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07-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC62422;
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                                                                                                                                                                                                                                                                                 reagents
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                                                                                                                                                                                                                                                                                                                                  The present sequence represents an uniform phosphodiester oligonucleotide. The specification describes oligomeric compounds containing 2.0-modified ribosyl nucleosides. The 2'-O-modified nucleosides include ring structures that position the sugar molety of the nucleosides preferentially in 3' endo geometries. The modified oligomeric compounds have increased binding affinity and increased nuclease resistance. The oligomeric compounds can be used in diagnostic, therapeutic and research reagents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= b
/note= "2'-O-[2-N,N-dimethylaminoethyl)oxyethyl-5- methyl
uridine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligomeric compound; 2'-O-modified ribosyl nucleoside; 3' endo geometry; nuclease resistance; phosphorothioate; ss.
                                                                                                                                                                                                                                         Novel 2'-O-aminoethyloxyethyl modified nucleosides and oligonucleotides used in diagnostic, therapeutic and research reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag= a
'note= "nucleosides linked by phosphodiester linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2'-O-modified ribosyl oligonucleotide with phosphodiester linkages.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 19; 7.7e+02; hes 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 19; DB
100.0%; Pred. No. 7.7
ative 0; Mismatches
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                                                                                                                                                                                                                                                                                                       Disclosure; Page 44; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2709 AAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAAA
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                                                                                           98US-00130566
                                                          99WO-US017895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC
                                                                                                                                  (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                       Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                          WPI; 2000-205668/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-1998;
                                                                                                                                                                       Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Manoharan M,
                                                          06-AUG-1999;
                                                                                               07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-FEB-2000
                     17-FEB-2000
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The present invention is concerned with methods of determining the nuclease stability of oligomeric compounds using capillary-gel electrophoresis and laser-induced fluorescence. The methods are useful in the polymerase chain reaction (PCR), molecular cloning and disease diagnosis and treatment. The present sequence was used in a demonstration of the methods of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                   The present sequence represents an oligomeric compound containing 2'-0-modified ribosyl nucleosides. The oligomeric compound contains phosphodiester linkages. The 2'-0-modified nucleosides include ring structures that position the sugar moiety of the nucleosides preferentially in 3' endo geomeries. The modified oligomeric compounds have increased binding affinity and increased nuclease resistance. The oligomeric compounds can be used in diagnostic, therapeutic and research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Determining the nuclease stability and relative binding affinity of an oligomeric compound comprises capillary gel electrophoresis using laser-induced fluorescence.
                                                       Novel 2'-O-aminoethyloxyethyl modified nucleosides and oligonucleotides used in diagnostic, therapeutic and research reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T19 diester; nuclease stability assay; polymerase chain reaction; PCR; molecular cloning; disease diagnosis; disease treatment; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 19; DB 1; Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                119 diester for use in nuclease stability assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Col 19-20; 14pp; English.
                                                                                                                                                                               Disclosure; Page 51; 60pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cummins LL;
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WPI; 2000-205668/18
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Best Local Similarity
Matches 19; Conserv
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AAZ95240/
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                                                                                                                                                                                                                                                                                                          //note= "Optionally all 3'-0-(2-methoxyhexyl) or all 2'-0-
(2-methoxyethyl)"
                                                                                                                                                                  phosphorothioate; gene therapy; ISIS # 22111;
                                                                                                                                                                                                                                                        *tag= d
note= "Optionally all phosphorothioate internucleotide
                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel modified oligonucleotides, useful in antisense methodologies, diagnostics, therapeutics and as research reagents.
                                                                                                                                                                                                                                *tag= a
note= "Phosphorothioate internucleotide linkage"
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Pred. No. 7.7e+02;
               Indels
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      Pred. No. 7.7e+02; i Mismatches 0;
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100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                                Modified oligonucleotide #3 ISIS # 22111.
                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 54; Page 59; 75pp; English.
                                                                                                                                                                           research reagent; therapeutic; ss
                              2709 AAAAAAAAAAAAAAAAA 2727
100.0%; Pr. 0;
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/mod_base= OTHER
                                                 AAAAAAAAAAAAAA
                                                                                            BP
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AAZ95241 standard; DNA; 19
                                                                                                                                                                  Antisense oligonucleotide;
                                                                                                                              (first entry)
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Best Local Similarity 100.0
Matches 19, Conservative
                                                                                                                                                                                                                                                                           inkages"
               Conservative
                                                                                                                                                                                                                                               15. .19
                                                                                                                                                                                                                                                                                     6. .19
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                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-182445/16.
      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                       WO200004189-A1
                                                                                                                                                                                                                                                                                   modified base
                                                                                                                              05-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                               Manoharan M,
                                                                                                                                                                                                                      misc_feature
                                                                                                                                                                                                                                                misc_feature
                                                                                                                                                                                                                                                                                                                                                                                          13-JUL-1999;
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               19;
                                                                                                                                                                                            Synthetic.
                                                19
                                                                                                             AAZ95241;
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              Matches
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This sequence represents a modified oligonucleotide used in the course of the invention. The invention relates to oligonucleotides comprising nucleotides covalently linked together by internucleotide linkages where at least 1 nucleotide is linked to adjacent nucleotide by a 2',5'- internucleotide linkage and bears a 3'-substituent. The oligonucleotides can be used in gene therapy and are also useful in antisense methodologies, diagnostics, therapeutics and as research reagents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "Optionally all 3'-O-(2-methoxyhexyl) or all 2'-O-
(2-methoxyethyl)"
                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide; phosphorothioate; gene therapy; ISIS # 22110; research reagent; therapeutic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= d
note= "Optionally all phosphorothioate internucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel modified oligonucleotides, useful in antisense methodologies, diagnostics, therapeutics and as research reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/note= "Phosphorothioate internucleotide linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                  Modified oligonucleotide #3 ISIS # 22110,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 54; Page 59; 75pp; English.
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19 ААААААААААААААА
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                                                                                                                          AAZ95240 standard; DNA; 19
                                                                                                                                                                                                                                (first entry)
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200004189-A1
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                                                                                                                                                                                                                                05-JUN-2000
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                                                                                                                                                                             AAZ95240;
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RESULT 771 AAA06839/c

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Gaps

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Oligonucleotide; nuclease resistance; psoriasis; antipsoriatic; dermatological; cytostatic; virucide; antibacterial; fungicide; therapy; diagnosis; DNA-RNA hybrid; ss.
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                                                                    Oligonucleotide ISIS 22115
                                  05-MAR-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-NOV-2000
                                                                                                                                                                           Synthetic
 AAA88952;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          misc_RNA
                                                                                                                                                                                                             Key
The invention relates to aminooxy-modified nucleosides and oligonucleotides and to oligonucleotides that elicit RNase H for cleavage in a complementary nucleic acid strand. It also relates to oligonucleotides wherein at least some of the nucleotides are functionalised to be nucleas resistant, at least some of the nucleotides are include a substituent that potentiates hybridisation of the coligonucleotide to a complementary strand, and at least some of the nucleotides include a 2'deoxy-erythro-pentofuranosyl sugar moiety. The nucleotides include a 2'deoxy-erythro-pentofuranosyl sugar moiety. The provides for improved binding of such oligonucleotides to a complementary strand. The oligonucleotides of the invention are used as diagnostic, therapeutic or research reagents, and can be used to modulate gene cypression in organisms. The oligonucleotides containing the modified nucleosides have increased nuclease resistance and increased binding affinity to a complementary strand. The present sequence represents an oligonucleotide containing nucleotides substituted with a 2'-O-{2- [N-(2- amino)ethyl-N-(methyl)]aminooxyethyl} group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aminooxy-modified nucleosides and oligonucleotides useful in diagnostic, therapeutic and research reagents and for modulating the expression of
                                                                                                                                                                                                                                                                                  /*tag= a
/note= "These nucleotides are substituted with 2'-O-{2-
[N-(2-amino)ethyl-N-(methyl)]aminooxyethyl} group"
                                                                                                                                         Modified nucleoside; aminooxy group;
2'-deoxy-erythro-pentofuranosyl sugar moiety; nuclease resistant;
hybridisation; binding affinity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
lve 0; Mismatches 0; Indels
                                                                                                        Modified T-containing oligonucleotide, SEQ ID NO:14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kawasaki AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 99; Page 120; 195pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prakash TP,
                                                                                                                                                                                                                                        Location/Qualifiers
   AAA06839 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US017988
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                                                                        (first entry)
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Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein in organisms.
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                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-1998;
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                                                                        19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                       17-FEB-2000
                                                                                                                                                                                                               Synthetic
                                     AAA06839
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/\*tag= a /mod\_base= OTHER /note= "2'-0-(2-methoxyethyl)thymidine"

"phosphorothioate linkage"

note=

\*tag=

location/Qualifiers

/mod\_base= OTHER /note= "2'-0-(2-methoxyethy1)thymidine"

Ω

\*tag=

/mod\_base= OTHER /note= "2'-0-(2-methoxyethy1)thymidine"

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/\*tag=

/\*tag= e /label= RNA

/mod\_base= OTHER /note= "2'-O-(2-methoxyethyl)uridine"

U

/\*tag=

99US-00303586.

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Oligonuclectide ISIS 22115 contains a mixed phosphodiester and phosphorothioate backbone and has 2'-O-(2-methoxyethyl) chemistry. It was used in experiments to determine the effects of snake venome phosphodiesterase and liver homogenate on the stability of and liver homogenate on the stability of and B-form conformational geometry. The A-form geometry and outless the binding affinity and nuclease resistance of the oligonuclectide. The B-form geometry allows the oligonuclectide to serve as substrate for RNase-H when bound to a target nucleic acid strand. The oligonuclectides can be used to treat psoriasis and other inflammatory skin conditions, skin clancers and viral, bacterial and fungal infections, and in various
                                                                                                                                                       to
                                                                                                                                           New oligonuclectides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
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Pred. No. 7.7e+02;
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Mohan V;
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Best Local Similarity
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2709 AAAAAAAAAAAAAAAA 2727

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19

100.08;

AAA88952 standard; DNA; 19 BP

RESULT 772 AAA88952,

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WO200066609-A1
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modified_base
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                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                AAA88949;
                                                     Query Match
Best Local
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                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents 2'-modified chimeric oligonucleotides containing "-modified T. The nucleotides were used to examine the effects of the modifications on nuclease resistance. Novel oligonucleotides of the invention have both A- and B-form conformational geometry. The A-form geometry modulates the binding affinity and nuclease resistance of the oligonucleotide. The B-form geometry allows the oligonucleotide to serve as substrate for RNase-H when bound to a target nucleic acid strand. The oligonucleotides can be used to treat psoriasis and other inflammatory
                                                                                                                                                                    Oligonucleotide; nuclease resistance; psoriasis; antipsoriatic; dermatological; cytostatic; virucide; antibacterial; fungicide; therapy; diagnosis; ss.
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C
                                                                                                                                                                                                                                                                                                                                                                                                /*tag= d
/mod_base= OTHER
/note= "2'-modified thymidine, i.e. -S-Me, -Me, 2'-ara-
(F), 2'-ara-(OH), -2'-ara-(OMe)"
                                                                                                                                                                                                                                                                       2'-ara-
                                                                                                                                                                                                                                                                                                                                                                      -Me, 2'-ara-
                                                                                                                                                                                                                                                                                                                       -Me, 2'-ara-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New oligonucleotides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
       Gaps
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(F), 2'-ara-(OH), -2'-ara-(OMe)"
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/mod_base= OTHER
/mote= "2.-modified thymidine, i.e. -S-Me,
(F), 2'-ara-(OH), -2'-ara-(OMe)"
                                                                                                                                                                                                                                                                                                                                                 *tag= c
mod_base= OTHER
note= "2'-modified thymidine, i.e. -S-Me,
(F), 2'-ara-(OH), -2'-ara-(OMe)"
       Indels
       ..
     Mismatches

    Modified chimeric oligonucleotide.

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                        2709 AAAAAAAAAAAAAAAA 2727
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/mod base= OTHER
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                                          19 AAAAAAAAAAAAAAA 1
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AAA88965 standard; DNA; 19 BP.
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                                                                                                                                (first entry)
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200066609-A1
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modified_base
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                                                                                                                                                                                                            Synthetic
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    Matches
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skin conditions, skin cancers and viral, bacterial and fungal infections, and in various diagnostic applications
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotides containing sequences with A and B geometry, used to treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
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                                                                                                                                                                        Gaps
                                                                                                                                                                        ö
                                                                                                                        Length 19;
                                                                                                                                                                      0; Indels
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/mod_base= OTHER
/note= "3'-0-(2-methoxyethyl)thymidine"
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                                                                      Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      '*tag= e
'note= "phosphorothioate linkage"
                                                                                                                 ch 0.7%; Score 19; DB 1; I
1 Similarity 100.0%; Pred. No. 7.7e+02;
19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide; nuclease resistance; dermatological; cytostatic; virucide;
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AAA88949 standard; DNA; 19
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*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MPI; 2000-672833/65.
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RESULT 776
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the effects of snake venom phosphodiesterase and liver homogenate on the stability of oligonucleotides. Novel oligonucleotides of the invention have both A- and B-form conformational geometry. The A-form geometry modulates the binding affinity and nuclease resistance of the oligonucleotide. The B-form geometry allows the oligonucleotide to serve as substrate for RNase-H when bound to a target nucleic acid strand. The oligonucleotides can be used to treat psoriasis and other inflammatory skin conditions, skin cancers and viral, bacterial and fungal infections, and in various diagnostic applications
                                                                                                                                                                                                                                                                          Oligonucleotide; nuclease resistance; psoriasis; antipsoriatic; dermatological; cytostatic; virucide; antibacterial; fungicide; therapy; diagnosis; DNA-RNA hybrid; ss.
                                                                                                                            Gaps
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0
                                                                                                          Length 19;
                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                            note= "2'-0-(2-methoxyethy1)thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                              note = "2'-0-(2-methoxyethyl)thymidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "2'-0-(2-methoxyethyl)uridine"
                                                                                        Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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/*tag= f
/note= "phosphorothioate linkage"
                                                                                                         DB 1; Le
                                                                                                        0.7%; Score 19; DB
100.0%; Pred. No. 7.7
:ive 0; Mismatches
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mod base= OTHER
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/mod_base= OTHER
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/mod_base= OTHER
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label= RNA
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                                                                                                                                                                                                                                                            Oligonucleotide ISIS 22113
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                                                                                                                  Local Similarity 100 nes 19; Conservative
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                                                                                                           Query Match
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Matches
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                                            New oligonucleotides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
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100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "3'-0-(2-methoxyethyl)thymidine"
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/note= "3'-O-(2-methoxyethy1)thymidine"
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/note= "3'-O-(2-methoxyethy1)thymidine"
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/note= "3'-0-(2-methoxyethy1)thymidine"
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"phosphorothioate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and in various diagnostic applications
                                                                                                                                                            Example 54; Page 69; 132pp; English.
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Best Local Similarity luv...
Loca 19; Conservative
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WPI; 2000-672833/65.
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                                                                                            to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide, nuclease resistance; psoriasis; antipsoriatic; dermatological; cytostatic; virucide; antibacterial; fungicide; therapy; diagnosis; ss.
                                                                                        New oligonucleotides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                      Oligonucleotide ISIS 22114 contains a mixed phosphodiester and
                                                                                                                                                                                                                                                                                                            0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
Ative 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "3'-0-(2-methoxyethyl)thymidine"
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                                                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                   Example 54; Page 69; 132pp; English
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/mod_base= OTHER
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           99US-00303586,
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Matches 19; Conservative
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                             (ISIS-) ISIS PHARM INC
                                                  Mohan V;
                                                                      WPI; 2000-672833/65.
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modified_base
                                                  Manoharan M,
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          03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
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100.0%; Pred. No. 7.7e+02;
cive 0; Mismatches 0; Indels
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/mod_base= OTHER
/note= "2'-0-(2-methoxyethyl)thymidine"
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/note= "2'-0-(2-methoxyethy1)thymidine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and in various diagnostic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 54; Page 69; 132pp; English.
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/mod base=
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WO200066609-A1.
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Cole DL;

Krotz A,

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This invention describes a novel synthetic method (M) comprising: (a) providing a sample comprising a number of oligomers of formula (1); (b) contacting the sample with a deprotecting agent to remove R t groups from the oligomers; and (c) reacting the oligomer with a cleaving reagent. The method is used to produce oligomeric compounds for use in antisense and oligonucleotide therapies. The method enables the synthesis of oligomers with a reduction in the number acrylonitrile groups attached.

Acrylonitrile has been demonstrated to be a potent carcinogen in rats. This sequence represents a phosphorothioate 20-mer primer which is used in the method of the invention
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                                                                                                                                                                                                                                           Novel method for the production of oligomers with reduced exocyclic adducts comprises treatment with deprotecting and cleaving reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid cleavage; solid support; DNA-RNA hybrid;
affinity chromatography; sequencing; mutagenesis; DNA preparation;
nucleic acid purification; ss.
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Pred. No. 7.7e+02;
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               99EP-00307066
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                                                                                                                                                  Manoharan M,
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Matches 19; Conservative
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                                                                                                                                                  Ravikumar VT,
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               06-SEP-1999;
                                                     04-FEB-1999;
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misc_RNA
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AAC62454/
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                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotides containing sequences with A and B geometry, used treat and diagnose e.g. psoriasis, skin cancers and viral, fungal and bacterial infections, bind to single stranded RNA or DNA.
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                                                                                              /note= "2'-0-(2-methoxyethyl)uridine"
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/note= "phosphorothioate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and in various diagnostic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Phosphorothioate 20-mer primer DNA #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 54; Page 69; 132pp; English.
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/mod_base= OTHER
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/label= RNA
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/*tag= a
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                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                            Manoharan M, Mohan V;
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-672833/65
                                                                                                                                      WO200066609-A1
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modified_base
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                                                                                                                                                                         09-NOV-2000
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Best Loca Matches

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RESULT 779 AAA71630/

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Gaps

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Length 19; 0; Indels

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19 AAAAAAAAAAAAAAA 1
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                                                          RESULT 782
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ID AAH4
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incorporated at predetermined site from a solid support involves cleaving the nucleic acid molecule at the site of unconventional nucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to nucleotide oligomers comprising monomer units. Oligomers modulate gene expression when hybridized by a single- or double-stranded nucleic acid. They are useful for gene therapy, diagnostic and investigative purposes
                                                                                   The present invention is concerned with the cleavage of nucleic acids from solid supports. This is carried out by adding a non-conventional nucleotide into the nucleic acid attached to the support, so that it is recognised and cleaved by a specific DNA glycosylase and the sequence is sequencing, in vitro amplifications, cDNA and template preparation, DNA-based assays, mutagenesis procedures, nucleic acid purification and affinity chromatography. The present sequence is an oligonucleotide used in assays to demonstrate the methods of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Guanidinium functionalized oligomers prepared from corresponding monomer units, are hybridizable with a specific RNA or DNA sequence, useful for diagnostic and therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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0.7%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 7.7e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mohan V;
                                                                                                                                                                                                                                                                                                            100.0%; Preu. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gene expression; gene therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prakash TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 26; Page 54; 108pp; English
                                                    Example 3; Page 34; 47pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF31458 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                               07-JUL-2000; 2000WO-US018609
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide ISIS 109989
                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100."
Matches 19, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF31458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 781
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2709 AAAAAAAAAAAAAAAA 2727

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New oligomers for use as research reagent, for treating disease caused by undesired production of proteins, and for diagnosing and treating AIDS, atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to C3' methylene hydrogen phosphate oligomers. The oligomers may be used as research reagents, for treating disease caused by undesired production of proteins and for diagnosing and treating AIDS and atherosclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                  oligomer; C3' methylene hydrogen phosphate; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 0 G; 15 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 19; DB 1; L. Pred. No. 7.7e+02;
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100.0%; Pred. No. ...
0; Mismatches
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/mod_base= OTHER
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AAF31564 standard; DNA; 19
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                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                          Cook PD, Manoharan M,
                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-138117/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                    ISIS sequence 32327.
                                                                                                                                  DNA/RNA hybrid; oligatherosclerosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide #8.
                                                                                                                                                                                                                   WO200102419-A1.
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modified_base
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                                                                  09-APR-2001
                                                                                                                                                                                                                                                      11-JAN-2001.
                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                 Preparing sulfurized 2' substituted phosphorothioate oligonucleotides useful in biological research, comprises phosphitylating the 5'-hydroxyl of a nucleic acid having a nucleoside with a 2' modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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/note= "optionally 3'-O-(2-methoxyethyl) or 2'-O-(2-
methoxyethyl)"
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/note= "optionally phosphorothioate backbone"
16. .19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 19; 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                      /note= "Modified with 2'-0-methoxyethyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; RNase H type II; RNase H1 cleavage substrate; gene therapy; primer; phosphorothioate backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human type II RNase H substrate oligonucleotide #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                             Cheruvallath ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 19;
100.0%; Pred. No.
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           /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                             Example 12; Col 7; 7pp; English
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                                                                                                                          11-JAN-2000; 2000US-00481486
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/*tag= a
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Matches 19, Conservative
                                                                                                                                                                                                             Ravikumar VT,
                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
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modified_base
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                                                                    US6242591-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, RNase H type II, RNase H1 cleavage substrate; antisense therapy;
gene therapy; primer; phosphorothioate backbone; ss.
                                                                                                                                                                  Chimeric oligonucleotides that can serve as substrates for human RNase H1, useful for enhancing the effectiveness of antisense gene therapies.
                                                                                                                                                                                                                                                                                                                                                                                    Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "optionally 3'-0-(2-methoxyethyl) or 2'-0-(2-
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/note= "optionally phosphorothioate backbone"
16. .19
                                                                                                                                                                                                                                                                                                                                                           Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human type II RNase H substrate oligonucleotide #5.
                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 19; DB 1; Lv
100.0%; Pred. No. 7.7e+02;
:ive 0; Mismatches 0;
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                                                                                                                      Manoharan
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                            methoxyethyl)"
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                                             29-SEP-2000; 2000WO-US026729
                                                                       99US-00409926
                                                                                                                      Wu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH25738 standard; DNA; 19
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Best Local Similarity 100.
Matches 19; Conservative
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                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                      Lima WF,
                                                                                                                                              WPI; 2001-343164/36.
WO200123613-A1
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                                                                       30-SEP-1999;
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                       05-APR-2001
                                                                                                                        Crooke ST,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
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Manoharan M;

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can act as substrates for human RNase HI (a type II RNase). The sequence consists of two portions, one of which is capable of supporting cleavage of a complementary target RNA and the other of which is incapable of supporting such cleavage. These can be used to enhance the effectiveness of antisense therapies. The present sequence is an RNase H substrate used in the exemplification of the invention
                                                                                                                                     The present invention provides a number of DNA-RNA oligonucleotides which
                                                                            Chimeric oligonucleotides that can serve as substrates for human RNase {\rm HI}_{\rm c} useful for enhancing the effectiveness of antisense gene therapies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New 2'-O-acetamido modified nucleosides (I) used to produce oligonucleotides which have enhanced nuclease resistance and superior hybridization properties than prior art.
                                                                                                                                                                                                                               Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fraser AS, Prakash TP,
                                                                                                              Example 54; Page 88; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00378568.
                                 Wu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00378568
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          (ISIS-) ISIS PHARM INC
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Best Local Similarity
                                Lima WF,
                                                       WPI; 2001-343164/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
modified_base
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                                                                                                                                                                                                                                                                              19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-NOV-2000
                                 Crooke ST,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                         AAC83664;
                                                                                                                                                                                                                                                                              Matches
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                                  Gaps
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0
   Length 19;
                                 0; Indels
0.7%; Score 19; DB 1; Le
100.0%; Pred. No. 7.7e+02;
cive 0; Mismatches 0;
                                                           2709 AAAAAAAAAAAAAAAA 2727
                                                                                         19 AAAAAAAAAAAAAA 1
                                 Conservative
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2'-O-N-[2-(dimethylamino)ethylacetamido]-modified oligo ISIS #32335. 2'-O-acetamido; diagnostic; kinase modulator; nuclease resistance; AAC83664 standard; DNA; 19 BP 02-MAR-2001, (first entry)

/mod\_base= OTHER /note= "2'-0-N-[2-(dimethylamino)ethylacetamido]SMeU" tumour formation, cancer; protein kinase C expression; cell adhesion molecule expression; multidrug resistance; ss. Location/Qualifiers \*tag= a

Kawasaki AM;

Example 12; Col 28; 29pp; English.

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modified nucleosides were used to produce oligonaclectides which have enhanced nucleosides were used to produce oligonaclectides which have enhanced nucleosides were used to produce oligonation properties than prior art. The oligomeric compounds are useful for identification or quantification of ribonacleic acid and deoxyribonacleic acid or for modulating the activity of an ribonacleic acid or deoxyribonacleic acid modulating the activity of an ribonacleic acid or deoxyribonacleic acid molecule. They have a modified nucleoside monomer and are specifically hybridisable with a preselected nucleotide sequence of a single-stranded or double-stranded target deoxyribonacleic acid or ribonacleic acid molecule. The oligomers are further useful in a ras-luciferase fusion system using ras-luciferase transactivation. They are useful in abnormal cell proliferation and tumour formation and modulation of expression of protein kinase C and cell adhesion molecules such as ICAM. They are useful in the modulation of proteins related to multidrug resistance and curral genomic nucleic acids such as HOV, herpes viruses, Epstein-Barr virus, cytomegalovirus, papillomavirus, hepatitis C virus and influenza
present sequence is a modified oligonucleotide. 2'-0-acetamido-
88888888888888888888888888888888888
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Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö Score 19; DB 1; Length 19; Pred. No. 7.7e+02; 0; Indels 100.0%; Prec. ... 0.7%; 19; Conservative Query Match Best Local Similarity Matches

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2709 AAAAAAAAAAAAAAA 2727 19 AAAAAAAAAAAAAAA 1 요 ð

Βb AAK98526 standard; DNA; 19 (first entry) 16-APR-2002 AAK98526; RESULT 787 AAK98526/

Target detection, quantitative analysis; probe, medical diagnosis; forensics; bacterial screening; tissue typing; gene expression analysis; Nucleic acid quantitative analysis related oligonucleotide #1.

Location/Qualifiers genotyping; ss. Synthetic. 

modified\_base

/mod\_base= OTHER /note= "modified by thiol" /\*tag=

WO200202810-A2

10-JAN-2002

02-JUL-2001; 2001WO-EP007575

01-JUL-2000; 2000DE-0103334.

(CLON-) CLONDIAG CHIP TECHNOLOGIES GMBH.

Kaiser Ermantraut E, Ellinger T, Ehricht R, Wagner G; Bickel R, Schulz T,

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WPI; 2002-154760/20

Determining targets by interaction with probe array, useful e.g. for diagnosis, based on detecting formation of precipitate at specific probe

Page 47; 92pp; German. ŝ Example

The present invention relates to a method for the qualitative and

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alkylation of alcohols, amines, or thiols, useful for preparing nucleosides that are precursors for preparation of oligomeric compounds beneficial as therapeutics, involves use of cyclic sulfate intermediates.
quantitative detection of targets in a sample by molecular interaction between the target and probes in an array. The method can be used to detect interactions between nucleic acids, antigens and antibodies or receptor and ligands, particularly in applications such as medical diagnosis, forenatic schence, bacterial screening, tissue typing for transplantation, monitoring gene expression, and genetying. The present sequence is a modifying oligonucleotide used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                     2'-0-alkyl oligonucleotide; nuclease resistance; diagnosis; therapy; ss.
                                                                                                                                                                   Gaps
                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kawasaki AM;
                                                                                                                                      Score 19; DB 1; Length 19; Pred. No. 7.7e+02; 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= b
/mod_base= OTHER
/note= "2'-methyl thioethyl thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= c
/mod_base= OTHER
/note= "2'-methyl thioethyl thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thioethyl thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thioethyl thymidine'
                                                                                                                 Seguence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jung ME,
                                                                                                                                                                                                                                                                                                                                                              Methyl thioethyl modified oligonucleotide
                                                                                                                                         0.7%; Score 19;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
/mod_base= OTHER
/note= "2'-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "2'-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Col 35; 45pp; English
                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                  ВР
                                                                                                                                                     ilarity 100.0%;
Conservative 0
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                                                                                                                                                     Local Similarity
les 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_baве
                                                                                         the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                           ABA91949;
                                                                                                                                          Query Match
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                                                                                                                                                         Best Loca
Matches
                                                                                                                                                                                                                                                          RESULT 788
                                                                                                                                                                                                                                                                     ABA91949
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The present sequence is that of a chimeric oligonucleotide having some

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cc with methoxyethoxy (see ABA91950) and dimethylaminopropyl (see ABA91951)

with methoxyethoxy (see ABA91950) and dimethylaminopropyl (see ABA91951)

cc with methoxyethoxy (see ABA91950) and dimethylaminopropyl (see ABA91951)

cc revealed the nuclease resistance of the modified oligomers. The invention of provides methods for the alkylation of alcohols, amines, thiols and their cc derivatives by cyclic sulfate intermediates. In particular, methods for the alkylation of the 2', 3' or 5'-balkyl cc the alkylation of the 2', 3' or 5'-balkyl culfate modified compounds are disclosed. Displacement of the 2', 3' or 5'-O-alkyl cc sulfate with a nucleophile provides 2', 3' or 5'-O-modified compounds are disclosed. Displacement of the 2', 3' or 5'-O-alkyl nucleosides and nucleosides and nucleoside cc the preparation of 2'-O-alkyl nucleosides, nucleosides and nucleoside cc surrogates that are precursors for the preparation of oligomeric compounds useful as therapeutics, diagnostics and research reagents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2'-0-alkyl oligonucleotide; nuclease resistance; diagnosis; therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   *tag= a
|mod_base= OTHER
'note= "2'-dimethylaminopropyl thymidine"
                                                                                                                                                                                                                                             Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note= "2'-dimethylaminopropyl thymidine"
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/note= "2'-dimethylaminopropyl thymidine"
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                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                            Score 19; DB 1; L
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dimethylaminopropyl modified oligonucleotide.
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                                                                                                                                                                                                                                                      100.0%; Preα. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    БО,
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                                                                                                                                                                                                                                                                                                                                                                                                      ABA91951 standard; DNA; 19
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nes 19; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA91951;
                                                                                                                                                                                                                                                 Query Match
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Matches
          %$$$$$$$$$$$$$$$$$$$
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The present sequence is that of a chimeric compared with clienthylaminopropyl modifications. This was compared with coligonucleotides with methyl thioethyl (see ABA91949) and methoxyethoxy (see ABA91949) and methoxyethoxy (see ABA91949) and modifications for resistance to snake venom phosphodiesterase. The assay revealed the muclease resistance of the action of modified oligomers. The invention provides methods for the alkylation of aloohols, amines, thiols and their derivatives by cyclic sulfate controls, amines, thiols and their derivatives by cyclic sulfate in the compounds and their mediates. In particular, methods for the alkylation of the 2', 3' or 5'-0-alkyl sulfate modified compounds are strongers to farso of the 2', 3' or 5'-0-alkyl sulfate with a nucleophile provides 2', 3' or 5'-0-alkyl sulfate with a nucleophile provides 2', 3' or 5'-0-alkyl for the preparation of 2'-0-alkyl nucleotides, nucleosides and nucleoside surrogates that are precursors for the preparation of oligomeric compounds useful as therapeutics, diagnostics and research reagents
                                                                                       present sequence is that of a chimeric oligonucleotide having some 2'
nucleosides that are precursors for preparation of oligomeric compounds beneficial as therapeutics, involves use of cyclic sulfate intermediates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2'-0-alkyl oligonucleotide; nuclease resistance; diagnosis; therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "2'-methoxyethoxy thymidine"
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/mod_base= OTHER
/note= "2'-methoxyethoxy thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= b
/mod_base= OTHER
/note= "2'-methoxyethoxy thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "2'-methoxyethoxy thymidine"
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Methoxyethoxy modified oligonucleotide.
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                                                   Example 15; Col 35; 45pp; English.
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mod_base= OTHER
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Best Local Similarity 100.
Matches 19; Conservative
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modified_base
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The present sequence is that of a chimeric oligonucleotide having some 2'—methoxyethoxy modifications. This was compared with oligonucleotides with methyl thioethyl (see ABA91949) and dimethyl aminopropyl (see ABA91951) modifications for resistance to snake venom phosphodisterase. The always revealed the nuclease resistance of the modified oligomers. The invention provides methods for the alkylation of alcohols, amines, thiols and their derivatives by cyclic sulfate intermediates. In particular, methods for the alkylation of the 2', 3' or 5'-hydroxy position of concleosides and their analogues with cyclic sulfates to form the 2', 3' or 5'-o-alkyl sulfate modified compounds are disclosed. Displacement of the 2', 3' or 5'-o-alkyl sulfate with a nucleophile provides 2', 3' or 5'-o-alkyl sulfate with an ancleophile provides 2', 3' or 5'-o-alkyl sulfate with an ancleophile provides 2', 3' or 5'-o-alkyl sulfate with a nucleophile provides are especially useful for the preparation of 2'-o-alkyl nucleotides, nucleosides and their analogues. The methods are especially useful for the preparation of 2'-o-alkyl nucleosides surrogates that are precursors for the preparation of 2'-o-alkyl nucleosides and research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                   Alkylation of alcohols, amines, or thiols, useful for preparing nucleosides that are precursors for preparation of oligomeric compounds beneficial as therapeutics, involves use of cyclic sulfate intermediates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tailing reaction related exemplary primer biotin-dT18U SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tailing reaction, tailed primer, primer, probe, identification, detection, linear amplification scheme, chain extending enzyme,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                  Kawasaki AM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                  Jung ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Scor.
100.0%; Pred. No. '...
'... 0; Mismatches
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/mod_base= OTHER
/note= "biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                  Cook PD,
                                                                                                                                                                        Example 15; Col 35; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAA 2727
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25-MAY-2000; 2000US-00580358.
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                                  Fraser AS, Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19; Conservative
(ISIS-) ISIS PHARM INC.
                                                                   WPI; 2002-235143/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  telomerase; ss.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL51520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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ABL51520/c
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Gaps ; 0

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modified base
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                                                               Kawasaki AM,
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PART SERVICE S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                    The present invention describes a method for detecting a nucleic acid probe, which comprises using chain extending enzymes to elongate probes. The method comprises (a) treating the sample with a chain cerminating the sample with a chain necrminating the sample (b) contacting the sample with the probe containing a terminus capable of elongation by a chain extending enzyme, where the probe hybridises to the nucleic acid in the sample; (c) contacting the sample with a chain extending enzyme and its substrates, which elongates the probe; and (d) detecting the elongated hybridised probe. Also described is a method comprising; (a) treating nucleic acid molecules or modified nucleic acid in a sample with a reagent or reagents that render the nucleic acid sin a sample with a nucleic acid probe that includes an extendable terminus, under conditions where hybrids form; and (c) treating any hybrids formed with a non-template dependent chain elongating enzyme and its substrates, where any hybridised probe is extended. The method is useful for identifying and detecting nucleic acids, particularly DNA hybridisation probes. The present sequence the acids, particularly DNA hybridisation probes. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                       Identifying and detecting nucleic acids, particularly DNA hybridization probes, involves employing chain extending enzymes (e.g. telomerase) to elongate probes to render them readily detectable.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dihydroxy sugar moiety; 2'-O-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
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100.0%; Pred. No. 7.7e+02;
ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "2'-methoxyethoxy (MOE) residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 1 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    example from the present invention
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                                                                                                                                                                                                                                                     Example 1; Page 5; 10pp; English.
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/mod_base= OTHER
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                                                               Tullis RH, Streifel JA;
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(TULL) TULLIS R H. (STRE/) STREIFEL J A.
                                                                                                            WPI; 2002-361176/39.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-o-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of oligomeric compounds having improved hybridiastion affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dihydroxy sugar moiety, 2'-O-alkyl nucleotide, hybridisation affinity; nuclear resistance, alkylation, therapeutic; diagnostic; ss.
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                                                          Prakash TP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                                                             Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Manoharan M,
                                                             Manoharan M,
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                                                                                                                                                                                                                                                                                                                                                            Example 46; Col 31; 24pp; English.
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/*tag= a
/mod_base= OTHER
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Matches 19; Conservative
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                                                                Fraser AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
(ISIS-) ISIS PHARM INC
                                                                                                                          WPI; 2002-546338/58.
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Page 389

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Alkylating 2' position of 2',3'-dihydroxy sugar moiety of nucleoside used for preparation of 2'-O-alkylated compounds comprises dissolving nucleoside in aprotic solvent, cooling, treating with base, warming,
                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                     WPI; 2002-546338/58.
                                                                                                                                                                                                                                   08-JAN-1999;
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                                                                                                                                                                                                                                                            Kawasaki AM,
                                                                              Query Match
                                                                                                                      RESULT 794
                                                                                                                         AAD42004,
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The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in a least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with

Example 46; Col 33; 24pp; English.

cooling and reacting with ester.

Prakash TP;

Cook PD,

Manoharan M,

Fraser AS,

99US-00227782 99US-00227782

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                                                                                                                                                 The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-0-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dihydroxy sugar moiety; 2'-0-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
      cooling, treating with base, warming,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "5-methyl, 2'-dimethylaminooxyethyl residue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Prec. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19;
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                                                                                             Example 46; Col 33; 24pp; English.
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                                cooling and reacting with ester.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
/mod_base= OTHER
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nucleoside in aprotic solvent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%;
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Matches 19; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dipydroxy sugar moleties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-0-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide
a reactive ester. The method is useful for the preparation of 2'-O-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "2'-dimethylaminooxyethyl thymidine (T-2'DMAOE)"
18. .19
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dihydroxy sugar moiety, 2'-O-alkyl nucleotide, hybridisation affinity, nuclear resistance, alkylation, therapeutic, diagnostic, ss.
                                                                                                                                                                              Gaps
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                                                                                                                                           Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                                                                                                                                                             0; Indels
                                                                                                            Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook PD,
                                                                                                                                                  100.0%; Pred. wc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 46; Col 35; 24pp; English.
                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAA 2727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and reacting with ester.
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                                                                                                                                          0.7%;
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                                                                                                                                                                           19; Conservative
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                                                                                                                                          Query Match
Best Local Similarity
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modified_base
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                                                                                                                                                                                                                                                                                       RESULT 795
                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-0-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                             Dihydroxy sugar moiety, 2'-O-alkyl nucleotide, hybridisation affinity, nuclear resistance, alkylation, therapeutic, diagnostic, ss.
                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                 Oligonucleotide #23 used to illustrate the method of the invention.
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                                                            0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 19;
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                               Seguence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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larity 100.0%; Pred. No. 7.7e+02;
Conservative 0; Mismatches 0;
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used to illustrate the method of the invention
                                                                    100.0%; Pred. ....
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                            2709 AAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER
                                                                                                                                              AAD42020 standard; DNA; 19 BP.
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/mod_base=
                                                                                                                                                                                                                                                                                                     (first entry)
                                                                              Best Local Similarity 100.
Matches 19; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                                     04-NOV-2002
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                                                                                                                                                                                                                                                                      AAD42020;
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                                                                Query Match
                                                                                                                                                                                                         RESULT 796
AAD42020/c
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nucleoside used
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                                                                                                                                                                                                  Dihydroxy sugar moiety, 2'-O-alkyl nucleotide, hybridisation affinity;
nuclear resistance, alkylation, therapeutic, diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                     /*tag= a
/mod_base= OTHER
/note= "5-methyl, 2'-dimethylaminooxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alkylating 2' position of 2',3'-dihydroxy sugar moiety of nucleoside for preparation of 2'-0-alkylated compounds comprises dissolving nucleoside in aprotic solvent, cooling, treating with base, warming,
                                                                                                                                                                       Oligonucleotide #4 used to illustrate the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 19; DB 1; Lv
100.0%; Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 46; Col 31; 24pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cooling and reacting with ester.
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99US-00227782.
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                                                                                                                                               (first entry)
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                                                                                       DNA;
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                                                                                        AAD42001 standard;
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modified_base
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                                                                                                                                                                                                                                              Unidentified
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                                                                                                                                               04-NOV-2002
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AC AAD4
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Best Local Similarity Matches 19; Conserv

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                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-O-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of a ligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                 'note= "2'-dimethylaminooxyethyl thymidine (T-2'DMAOE)"
                                                  Dihydroxy sugar moiety; 2'-O-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dihydroxy sugar moiety; 2'-O-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
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                              Oligonucleotide #14 used to illustrate the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                              Cook PD,
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Pred. No. 7.7e+02;
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v 100.0%; Pred. No. ...
... 0; Mismatches
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                                                                                                        Location/Qualifiers
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/mod_base= OTHER
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ID AAD42005 standard; DNA; 19
           04-NOV-2002 (first entry)
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                                                                                                                  16. 19
                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
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Best Local Similarity
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modified_base
                                                                                   Unidentified
                                                                                                                                                                     US6403779-B1
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Alkylating 2' position of 2',3'-dihydroxy sugar moiety of nucleoside used for preparation of 2'-O-alkylated compounds comprises dissolving nucleoside in aprotic solvent, cooling, treating with base, warming, cooling and reacting with ester.
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                                                                                                         /note= "5-methyl, 2'-methoxyethyl residues"
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Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Cook PD,
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Location/Qualifiers
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                                                       /*tag= a
/mod base= OTHER
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/*tag=
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modified_base
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Kawasaki AM,
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WPI; 2002-546338/58.
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AAD41999/
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                                                                                                                 Alkylating 2' position of 2',3'-dihydroxy sugar moiety of nucleoside used for preparation of 2'-O-alkylated compounds comprises dissolving nucleoside in aprotic solvent, cooling, treating with base, warming,
                                                                                                                                                                                   The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with base, warming, cooling and reacting with nucleotides, nucleosides and nucleoside surrogates used for preparation of 1'-0-alkyl nuclear compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dihydroxy sugar moiety; 2'-O-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "5-methyl, 2'-aminooxyethoxy (2'-AOE) residues"
                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide #1 used to illustrate the method of the invention.
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                                                                            Prakash TP;
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100.0%; Pred. No. 7.7e+02;
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                                                                                                                                                                   Example 46; Col 33; 24pp; English
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                                                                                                                                                 cooling and reacting with ester.
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                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                             Kawasaki AM, Fraser AS,
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                                                          PHARM INC
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Best Local Similarity
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modified_base
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                     08-JAN-1999;
                                       08-JAN-1999;
                                                          SISI (-SISI)
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 11-JUN-2002
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Alkylating 2' position of 2',3'-dihydroxy sugar moiety of nucleoside used for preparation of 2'-O-alkylated compounds comprises dissolving nucleoside in aprotic solvent, cooling, treating with base, warming,
                                                                                                                                                                                                                                         The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-O-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dihydroxy sugar moiety, 2'-O-alkyl nucleotide, hybridisation affinity; nuclear resistance, alkylation, therapeutic; diagnostic; ss.
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/note= "5-methyl, 2'-dimethylaminooxyethoxy (2'-DMAOE)
residues"
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100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                  Example 46; Col 31; 24pp; English
                                                                                                                 cooling and reacting with ester.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19; Conservative
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                                               The present invention relates to a novel method of selective alkylation of the 2' position of 2', 3'-dihydroxy sugar moieties of a nucleoside. The method involves dissolving the nucleoside in at least one aprotic solvent, cooling, treating with base, warming, cooling and reacting with a reactive ester. The method is useful for the preparation of 2'-0-alkyl nucleotides, nucleosides and nucleoside surrogates used for preparation of of oligomeric compounds having improved hybridisation affinity and nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "2'-dimethylaminooxyethyl thymidine (T-2'DMAOE)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dihydroxy sugar moiety; 2'-0-alkyl nucleotide; hybridisation affinity; nuclear resistance; alkylation; therapeutic; diagnostic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
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31; 24pp; English.
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                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide; 2'-0-(2-(methylthio)ethyl)-5-methyluridine; antisense;
nuclear resistance, which are useful as therapeutics, diagnostics and research reagents. The present sequence is a modified oligonucleotide used to illustrate the method of the invention
                                                                                                              Gaps
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/mod_base= OTHER
/note= "2'-0-(2-methylthio)ethyl)-5-methyluridine"
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/note= "2'-O-(2-methylthio)ethyl)-5-methyluridine"
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                                                                                                                                                                                                                                                                                                                         Oligonucleotide with 2'-0-(2-(methylthio)ethyl)-5-methyluridine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Increasing binding of oligomeric compound to proteins useful in preparation of antisense therapeutics, involves use of modified oligomeric compound having oligonucleotide group.
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                                                                               0.7%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 7.7e+02;
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                                                     Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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/mod_base= OTHER
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28-JAN-2002; 2002US-00058740.
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                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                  DNA-RNA hybrid; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prakash TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                  ABZ58336;
                                                                                 Query Match
                                                                                               Local
                                                                                                                                                                                                            804
                                                                                                                                                                                                           RESULT 80.
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ВЪ.

ADE99265 standard; DNA; 19

ADE99265

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in comparison with the known 2'-O-(2-methoxyethyl) (2'-O-MOE) modification. The 2'-O-MTE modified oligonuclectides exhibited similar binding affinity to target RNA as their 2'-O-MOE equivalent while binding to human serum albumin was improved. The modification can be used to modulate the pharmacokinetics of oligonucleotides, e.g. in antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to oligomeric compounds having at least one nucleoside. The compounds are useful for therapeutic and investigative purposes and for treating hepatitis C virus infection. The compounds having 2'-0-modifications increases their affinity and nuclease resistance. This sequence represents an oligomeric compound of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           υ
                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligomeric compound having at least one nucleoside useful for therapeutic and investigative purposes \mathbf{e}.\mathbf{g}. for treating hepatitis
                                                                                                                                                                                                                                                                                                                                                                                    Oligomeric compound, hepatitis C virus, 2'-0-modification, nuclease resistance, hepatotropic, virucide, antiinflammatory; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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Pred. No. 7.7e+02;
                                                                                                                          DB 1; Length 19; 7.7e+02;
                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                Seguence 19 BP; 0 A; 0 C; 0 G; 15 T; 4 U; 0 Other;
                                                                                                                                                     ö
                                                                                                                0.7%; Scor.
100.0%; Pred. No. ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 5; 26pp; English.
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v 100.0%; Prev
0; P
                                                                                                                                                                               2709 AAAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2709 AAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                           Modified oligomeric compound #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                         AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                            ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-00370625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-00130566
                                                                                                                                                                                                                                                                            ADE99245 standard; DNA; 19
                                                                                                              Query Match
Best Local Similarity 10v..
Best Local 19; Conservative
                                                                                                                                                                                                                                                                                                                                12-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-895259/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6600032-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19
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                                                                                                                                                                                                                                                                                                       ADE99245;
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                                                                          therapy
                                                                                                                                                                                                                                                    RESULT 805
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Matches
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RESULT 806 ADE99265/c

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The invention relates to oligomeric compounds having at least one nucleoside. The compounds are useful for therapeutic and investigative purposes and for treating hepatitis C virus infection. The compounds having 2'-O-modifications increases their affinity and nuclease resistance. This sequence represents an oligomeric compound of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                 New oligomeric compound having at least one nucleoside useful for therapeutic and investigative purposes {\bf e}.{\bf g}. for treating hepatitis C
                                                                                            hepatitis C virus; 2'-0-modification;
hepatotropic; virucide; antiinflammatory; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nuclease resistance; hybrid binding; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/mod_base= OTHER
/note= "OTHER = 2'-O-[2-(guanidinium)ethyl]"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetically modified nuclease resistant oligomer #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 26; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2727
                                                                      Modified oligomeric compound #26.
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                                                                                                                                                                                                             99US-00370625.
                                                                                                                                                                                                                                   98US-00130566.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH97218 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                    Cook PD;
                                                                                               Oligomeric compound;
nuclease resistance;
                                                                                                                                                                                                                                                                                                            WPI; 2003-895259/82.
                                                                                                                                                                                                                                                                                                                                                               virus infection.
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modified base
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                                                                                                                                                            US6600032-B1
                                                                                                                                                                                                             06-AUG-1999;
                                                                                                                                                                                                                                    07-AUG-1998;
                                                 12-FEB-2004
                                                                                                                                                                                    29-JUL-2003
                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
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ADH97218/c
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Matches
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                                                                                                                                                   This invention relates to novel synthetically modified oligomers that have increased nuclease resistance and have enhanced hybrid binding. Such oligomers are useful for diagnostic and therapeutic uses such as antisense technologies. The invention also discloses a method for the preparation of the oligomers with modifications as fully defined in the specification. The present sequence represents a synthetically modified oligonucleotide of the invention.
                                                                                                   Guanidinium functionalized oligonucleotides used for diagnostic, therapeutic or investigative purposes comprises a number of nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Guanidinium functionalized oligonucleotides used for diagnostic, therapeutic or investigative purposes comprises a number of nucleotide
                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                     Nuclease resistance; hybrid binding; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note = "OTHER = 2'-0-[2-(guanidinium)ethyl]"
                                                                                                                                                                                                                                         Length 19;
                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                     Synthetically modified nuclease resistant oligomer #3
                                                                                                                                                                                                                        Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                        Score 19; DB 1; Le
Pred. No. 7.7e+02;
0; Mismatches 0;
                                                                    Mohan V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mohan V;
                                                                                                                                    Example 26; SEQ ID NO 7; Slpp; English
                                                                   Prakash TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                          100.0%; Fit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                base= OTHER
                                                                                                                                                                                                                                                                                         19 AAAAAAAAAAAAAAA 1
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                07-JUL-2000; 2000US-00612531.
                                 99US-00349040
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                                                                                                                                                                                                                                                          Conservative
                                                                  Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
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                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                   WPI; 2003-644179/61.
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                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-1999;
                                                                   Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Manoharan M,
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18-MAR-2003
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                                                                                                                                                                                                                                                         19;
                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                    ADH97214;
                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                  Local
                                                                                                                     units.
                                                                                                                                                                                                                                                                                                                   RESULT 808
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                                              This invention relates to novel synthetically modified oligomers that have increased nuclease resistance and have enhanced hybrid binding. Such oligomers are useful for diagnostic and therapeutic uses such as antisense technologies. The invention also discloses a method for the preparation of the oligomers with modifications as fully defined in the specification. The present sequence represents a synthetically modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to novel synthetically modified oligomers that have increased nuclease resistance and have enhanced hybrid binding. Such oligomers are useful for diagnostic and therapeutic uses such as an antisense technologies. The invention also discloses a method for the preparation of the oligomers with modifications as fully defined in the specification. The present sequence represents a synthetically modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guanidinium functionalized oligonucleotides used for diagnostic,
therapeutic or investigative purposes comprises a number of nucleotide
units.
                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nuclease resistance; hybrid binding; antisense technology; ss.
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                                                                                                                                                                                                                                                                                                     0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetically modified nuclease resistant oligomer #13
                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mohan V;
                                                                                                                                                                                                                                                                                                                       100.0%; Pred. ...
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26; SEQ ID NO 3; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                                                                  oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
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                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified base
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                                                                                                                                                                                                                                                                                                                                                       19;
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                                                                                                                                                                                                                                                                                                     Query Match
     Example
                                                                                                                                                                                                                                                                                                                                  Local
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Pred. No.

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Best Local Similarity 100.0%; P
Matches 19; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                    RESULT. 811
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes an oligonucleotide comprising several nucleotides covalently linked together by internucleotide linkages. At least one of the uncleotides is linked to an adjacent nucleotide by 2',5'. The uncleotide is linked to an adjacent nucleotide by 2',5'. The uncleotide linkage and bears a 3'-substituent. The oligonucleotides in thermucleotide linkage and hears a 3'-substituent. The oligonucleotides are useful; as antisense oligonucleotides; in pharmaceutical compositions of for treating organism having disease caused by undesired production of protein e.g. organism that utilises RNA-DNA transcription or RNA-Protein translation, bacteria, yeast, protozoa, algae and warm-blooded animals; of for developing diagnostic and therapeutic agents. The modified oligonucleotide exhibits improved properties of nuclease resistance and configuration of properties of nuclease resistance and exhibit configuration of action. This sequence represents an oligonucleotide of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide comprising at least one 2',5'-internucleotide linkage useful for treating organisms having disease caused by undesired production of protein e.g. bacteria, yeast, protozoa and algae.
                                                                                                                                                                                                                                                                                                                                                                                              antibacterial; protozoacide; antialgal; fungicide; internucleotide linkage; 3'-substituent; antisense; pharmaceutical; RNA-DNA transcription; RNA-protein translation; infection; diagnostic; therapeutic;
                                                                                                       Gaps
                                                                                                       ö
                                                                   Length 19;
                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                 Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                     0
                                                                   0.7%; Score 19; DB 1; Le
100.0%; Pred. No. 7.7e+02;
                                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                  Modified oligonucleotide seg id 6.
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 oligonucleotide of the invention.
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94WO-US010131.
96US-00602862.
98US-00115043.
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                                                                                                                                                                                                                                                             ADG28485 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nuclease resistance; ss
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                                                                                Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-1993;
02-SEP-1994;
28-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention.
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                                                                                                                                                                                                                                                                                              ADG28485;
                                                                     Query Match
                                                                                         Best Loca
Matches
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                                                                                                                                                                                                                                             ADG28485,
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DB 1; Length 19;

0.7%; Score 19;

Query Match

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                       Gaps
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/note= "2'-O-[2-(guanidinium)ethyl] thymidine"
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100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
                    Indels
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7.7e+02;
hes 0;
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                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                        BP.
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                                                                                                                                                                                                                          ADG47994 standard; DNA; 19
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/*tag= a
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(PRAK/) PRAKASH T P.
(MOHA/) MOHAN V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MANO/) MANOHARAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-031184/03
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modified_base
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ADG48004/c
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New oligomers containing guanidinium groups, useful for modulating gene expression by hybridizing oligomer with single- or double-stranded nucleic acids.
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        Oligonucleotide #5 used in the exemplification of the invention
                                                                                                                                                                                                                                                                                  /note= "2'-0-[2-(guanidinium)ethyl] thymidine"
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100.0%; Pred. No. 7.7e+02;
                                                            Hybridisation; diagnosis; therapeutic; investigation; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mohan V;
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mod base= OTHER
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18 19; Conservative
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(PRAK) PRAKASH T P.
(MOHA/) MOHAN V.
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modified_base
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modified_base
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                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligomers containing guanidinium groups, useful for modulating gene expression by hybridizing oligomer with single- or double-stranded nucleic acids.
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                                                                                                                                                             Oligonucleotide #11 used in the exemplification of the invention.
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/note= "2'-0-[2-(guanidinium)ethyl] thymidine"
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100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 26; SEQ ID NO 13; 54pp; English.
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                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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/mod_base= OTHER
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ADG48004 standard; DNA; 19
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                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (COOK/) COOK P D.
(PRAK/) PRAKASH T P.
(MOHA/) MOHAN V.
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Matches 19; Conserv
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Query Match

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WPI; 2004-118052/12
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                                                                                                                                                                                            The invention relates to a guanidinium functionalised nucleotide compounds. The guanidinium functionalised nucleotide compounds are used for preparation of oligomers useful for diagnostic, therapeutic and investigative applications. The 2-0-guanidinium ethyl modification increases binding affinity to a target. The present sequence represents a guanidinium functionalised oligonucleotide.
                                                                                                                                            New guanidinium functionalized nucleotide compounds useful for preparing oligomers used for diagnostic, therapeutic and investigative
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/note= "OTHER = 2-0-[2-(guanidinium)-ethyl] modified"
/mod_base= OTHER
/note= "OTHER = 2-0-[2-(guanidinium)-ethyl] modified"
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                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                      ss; guanidinium functionalised nucleotide; guanidinium; 2-0-guanidinium ethyl; increased binding affinity.
                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mohan V;
                                                                                                            Mohan V;
                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                              Example 26; SEQ ID NO 5; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cook PD, Prakash TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                             Prakash TP,
                                                                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                  100.08;
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                                                            99US-00349040
                                                                            99US-00349040
                                                                                                                                                                                                                                                                                                                                                                      ADH42931 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                           19; Conservative
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                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                             Cook PD,
                                                                                                                             WPI; 2004-118052/12.
                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 19; Conserv
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modified_base
                                                                                                                                                                applications.
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                                                                                                              Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JS6593466-B1
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                                                                             07-JUL-1999;
                                                            07-JUL-1999;
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                          US6593466-B1
                                           15-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
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New guanidinium functionalized nucleotide compounds useful for preparing oligomers used for diagnostic, therapeutic and investigative
                                                                                                                                                    The invention relates to a guanidinium functionalised nucleotide compounds. The guanidinium functionalised nucleotide compounds are used for preparation of oligomers useful for diagnostic, therapeutic and investigative applications. The 2-O-guanidinium ethyl modification increases binding affinity to a target. The present sequence represents guanidinium functionalised oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER = 2-0-[2-(guanidinium)-ethy]} modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2-0-[2-(guanidinium)-ethyl] modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Guanidinium functionalised oligonucleotide ISIS #109989.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; guanidinium functionalised nucleotide; guanidinium; 2-0-guanidinium ethyl; increased binding affinity.
                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mohan V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleotide c
therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 26; SEQ ID NO 4; 40pp; English.
                                                                                                               Example 26; SEQ ID NO 3; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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/mod_base= OTHER
/note= "OTHER =
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prakash
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-118052/12.
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modified_base
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The invention relates to 2'-O-aminoethylthioethyl-modified ribosyl nucleosides. The modified ribosyl nucleosides are used as monomers for the synthesis of modified antisense oligonucleotides, which are useful in diagnosis and therapeutics (e.g. in gene therapy, for treating organisms having a disease associated by the undesired production of proteins) and as research reagents. The oligonucleotides obtained from the monomers show enhanced hybrid binding affinity towards targeted DNA or RNA and resistance towards nucleases. This sequence represents a modified antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New 2'.-O-aminoethylthioethyl-modified ribosyl nucleosides useful ar
monomer for the synthesis of modified anti-sense oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                                                                                                                                                                                          2'-O-aminoethylthioethyl-modified ribosyl nucleoside;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 26; 26pp; English.
                                                                                                                                                                 Modified antisense oligonucleotide #25.
                                                                                                                                                                                                                            antisense oligonucleotide; ss.
                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-00370625.
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                                           ADJ77789 standard; DNA; 19
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                                                                                                                         (first entry)
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hes 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cook PD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Exemplary DNA molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-106293/11.
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                                                                                                                         06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                06-JAN-2004.
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                                                                                                                                                                                                                                                                    Synthetic.
                                                                                    ADJ77789;
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          RESULT 818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 819
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compounds. The guanidinium functionalised nucleotide compounds are used for preparation of oligomers useful for diagnostic, therapeutic and investigative applications. The 2-0-guanidinium ethyl modification increases binding affinity to a target. The present sequence represents guanidinium functionalised oligonucleotide.
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New 2'.-O-aminoethylthioethyl-modified ribosyl nucleosides useful a monomer for the synthesis of modified anti-sense oligonucleotides.
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                                                                                                                                                        0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2'-O-aminoethylthioethyl-modified ribosyl nucleoside;
antisense oligonucleotide; ss.
                                                                                                                       Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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Pred. No. 7.7e+02;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modified antisense oligonucleotide #5.
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99US-00370625
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                                                                                                                                                                                                    19; Conservative
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                                                                                                                                                                               Local Similarity
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Best Local Simi
Matches 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ77769;
                                                                                                                                                                                                                                                                              13
                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                          RESULT 817
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                                                                                                                                                                                                                                                     nanotube; nucleic acid sensor; DNA array; conductor; nanoparticle; biosensor; detection; screening; bacterial; viral; pharmaceutical; agricultural; food control; hygiene; environmental; forensic; nano-scale conductor; semiconductor; nano-electronic; prostatic nerve; bio-electronic interface; transistor; gated device; ss.
                                    Gaps
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                                   0; Indels
0.7%; Scor.
100.0%; Pred. No. ..
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The invention relates to nucleoside compounds. The nucleoside compounds are useful as antisense compounds in diagnostics, therapeutics, prophylaxis, and as research reagents and kits, and to prevent or delay infection, inflammation or tumour formation. The compounds have enhanced binding affinity properties. The present sequence represents a 2'-0-MOE-2-thio modified oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleoside compounds useful as antisense compounds to prevent or
                                                                                                                                                                                                                                               /note= "OTHER = 2'-0-[2-(methoxy-)ethyl]-2-thio-5-
methyluridine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 delay e.g. infection, inflammation or tumor formation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 19; DB 1; Lo
100.0%; Pred. No. 7.7e+02;
                                                ss; antisense; infection; inflammation; tumour; enhanced binding affinity.
2'-0-MOE-2-thio modified oligonucleotide #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. w.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rajeev KG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
16. 19
/*tag= a
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16. 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-AUG-2002; 2002US-00222588
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prakash TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MANO/) MANOHARAN M.
(PRAK/) PRAKASH T P.
(RAJE/) RAJEEV K G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-256363/24.
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                                                                                                                                                                             Key
modified_base
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                                                                                                                               Synthetic.
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ADO58963/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a nanotube (I) attached with one or more nucleic acid molecule(s). Also described: (1) chemically modifying (M1) a nanotubes; (2) physically modifying (M2) a nanotubes; (3) linking (M3) annotubes; (2) physically modifying (M2) a nanotubes; (3) linking (M3) annotubes; (4) a several linked nanotubes (II) produced by (M3); (5) directing (M4) nanotubes to specific targets; (6) a nucleic acid sensor (III) comprising (I), where the base sequence of the attached nucleic acid molecules being detected; (7) a DNA array consisting of an array of groups of one or more nucleic acid molecules being detected; (7) a DNA array consisting of an array of groups of one or more nucleic acid molecules of the same base sequence attached to each nanotubes in the group, and where the base sequence of the nucleic acid molecules, attached to the nanotubes in one group attached to acid molecules, attached to the nanotubes in one group and membrane support to which the DNA-molecules are attached; and a membrane support to which the DNA-modified nanotubes are attached; and conductor (IV) comprising (I). (I) is useful in coating one or more nanotubes with nanoparticles, which involves exposing (I) and a molecules, where the nanoparticles, which involves exposing (I) to comprising several attached complementary nucleic acid molecules, where the nanoparticles hybridise to the nucleic acid molecules, forming one or more coated nanotubes. (I) can be coher nanoparticles, forming one or more coated nanotubes. (I) can be coher nanoparticles, forming one or more coated nanotubes. (I) can be coher nanoparticles, in pharmaceutical applications, agricultural captured and control, hygiene and environmental monitoring and control, bygiene and environmental monitoring and control, applications, agricultural component in plant in a component in plant in a component in a component in plant in a component in a component in a component in the conductor or component in plant in a component in plant in a component in plant in a c
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                                                                                                                                                                                                                                                                                                                                                                         Carbon nanotube attached with one or more nucleic acid molecules, useful as biosensor for screening presence of bacterial or viral nucleic acid in clinical samples.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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100.0%; Pred. No. 7.7e+02;
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                                                                                                                                                                                                                               CSIR ) COMMONWEALTH SCI & IND RES ORG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 91; 147pp; English.
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ADM47150 standard; DNA; 19 BP.
                                                                                                                                    29-AUG-2003; 2003WO-AU001118
                                                                                                                                                                               30-AUG-2002; 2002AU-00951274
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                                                                                                                                                                                                                                                                                Mccall M, Moghaddam M;
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                                   WO2004020450-A1
                                                                                   11-MAR-2004
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ADM47150;

RESULT 820

ADM47150,

Query Match

Matches

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An isolated nucleic acid molecule, comprising nucleic acid sequence of
Nicotiana derived cytochrome P450 enzyme fragments, useful for producing
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                                                                                                                                                                                                                             enhancing renal uptake of an oligomeric compound. The method is useful for enhancing renal uptake of an oligomeric compound. The sequences of the invention are useful in diagnostics, therapeutics and as research reagents; and for treating infection caused by organisms (e.g. bacteria, yeast, protozoa and algae) in plants and higher animals. The present of 2'-O-modified oligomers. This sequence is used to illustrate enzymatic degradation method of the invention.
                                                                                                               Enhancing renal uptake of an oligomeric compound in the diagnostic and therapeutic applications involves incorporating at least one modified ribosyl nucleoside into the oligomeric compound.
                                                                                                                                                                                                                 The invention relates to 2'-0-modified ribosyl nucleosides and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; primer; PCR; cytochrome P450; transgenic; tobacco; plant.
                                                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
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                                                                                                                                                                                 Example 19; SEQ ID NO 5; 21pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tobacco cytochrome P450 PCR primer #6
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12-MAR-2002; 2002US-0363684P.
10-JAN-2003; 2003US-00340861.
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nes 19; Conservative
                                                    Cook PD;
     (MANO/) MANOHARAN M. (COOK/) COOK P D.
                                                                                 WPI; 2004-201317/19.
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                                                    Manoharan M,
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Matches
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                                                                                                                                                                                                                                                                                                                                                                       The invention relates to 2'-O-modified ribosyl nucleosides and methods of enhancing renal uptake of an oligomeric compound. The method is useful for enhancing renal uptake of an oligomeric compound. The sequences of the invention are useful in diagnostics, therapeutics and as research reagents; and for treating infection caused by organisms (e.g. bacteria, yeast, protozoa and algae) in plants and higher animals. The present sequence is an oligomucleotide used in animal studies. This sequence is used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                         Enhancing renal uptake of an oligomeric compound in the diagnostic and therapeutic applications involves incorporating at least one modified ribosyl nucleoside into the oligomeric compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to illustrate enzymatic degradation of 2'-0-modified oligomers.
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methoxyethyl or DMAEOE"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                          Example 19; SEQ ID NO 26; 21pp; English.
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                                                                                                             98US-00130566.
99US-00370625.
                                                                             06-FEB-2003; 2003US-00359328
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99US-00370625.
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Matches 19; Conservative
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                                                                                                                                                           (MANO/) MANOHARAN M. (COOK/) COOK P D.
                                                                                                                                                                                                                                            WPI; 2004-201317/19.
               US2004009938-A1.
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                                                                                                            07-AUG-1998;
06-AUG-1999;
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                                               15-JAN-2004
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transgenic tobacco plant, which involves operably linking (I) with a promoter functional in the plant to create a plant transformation vector, and transforming the plant with the plant transformation vector, selecting a plant cell transformed with the transformation vector, and regenerating a plant from the selected plant cell. The nucleic acid molecule is in an antisense orientation, sense orientation or is in a RNA interference orientation. The present sequence represents a PCR primer used to clone DNA encoding tobacco cytochrome P450 enzyme fragments of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic; anticonvulsant; notropic; muscula; anti-HIV; RNA interference; iRNA; antisense technology; lipid metabolism; cholesterol imbalance; dyslipidaemia hypercholesterolaemia; coronary attery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; disease; cancer; breast cancer; colon cancer; lung cancer; neurological disease; Huntington disease; spinocerebellar ataxia; viral disease; ADDS; hepatitis C virus; HCV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic;
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                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                                 Seguence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                               DB 1; Le
7.7e+02;
                                                                                                                                                                                               O.7%; Score 19; DB Local Similarity 100.0%; Pred. No. 7.7 tes 19; Conservative 0; Mismatches
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2003US-0454962P.
2003US-045594P.
2003US-0462894P.
2003US-046372P.
2003US-0465605P.
2003US-0465602P.
2003US-0469612P.
2003US-0469612P.
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2003US-0510246P.
2003US-0510318P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus.
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                                                                                                                                    the invention.
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25-APR-2003;
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The invention describes a RNA interference (IRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequences have one or more asymmetrical 2'-O alkyl modifications, the antisense cone or more asymmetrical phosphorothioate modifications and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (I); reducing (MI) apo8-100 cevels or glucose-6-phosphatase levels in a subject; producing (I); including (I), involves selecting a sequence with activity and introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (I) and instruction for its use; and adevice that can be dispense or administer a composition comprising (I). (I) is useful for reducing apo8-100 levels or glucose-6-phosphatase levels.

CC useful for reducing apo8-100 levels or glucose-6-phosphatase levels.

The subject is suffering from a disorder characterised by elevated or coherwise unwanted expression of apol-100, elevated or otherwise unwanted contains and or disersed cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from the HDL/LDL cholesterol imbalance,

CM sellidaemias, hypercholesterolaemia, statin-resistant

CM physicholesterolaemia, coronary artery disease (CMD), coronary heart disease (CMD) and atherosclerosis. (I) is administered to a subject to inhibit hepatic glucose production or for treating the diseases as mentioned above, cancer (e.g. breast, colon or treating the diseases as mentioned above, cancer (e.g. breast, colon or treating the diseases as mentioned above, cancer (e.g. breast, colon or spinocerbellar ataxia) or viral disease (e.g., Huntington or ligonucleotide that can creating the patitis C virus (RCV) antisense oligonucleotide that can control HCV gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic; anticonvulsant; nostropic; muscula; anti-HIV; RNA interference; iRNA; antisense technology; lipid metabolism; conclesterol imbalance; dyslipidaemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; glucose metabolism-related disorder; dasease; colon cancer; lung cancer; neurological disease; Huntington disease; spinocerebellar ataxia; viral disease; ADDS; hepatitis C virus; HCV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cardiant; vasotropic; antiarteriosclerotic; antidiabetic;
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sequence and antisense sequence which has specific modifications
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Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus (HCV) oligonucleotide seqid 6756.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                                                  Example 5; SEQ ID NO 6759; 378pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19; Conservative
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Best Local Similarity
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The invention describes a kWA interference (iKMA) agent (1) comprising a sense sequence and an antisense sequence, where the sense sequences have one or more asymmetrical phosphotochioate modifications come or more asymmetrical phosphotochioate modifications and the antisense sequence targets a human gene sequence. Also described and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (1); reducing (M1) apoB-100 (control one or more asymmetrical modification in the sequence, where the modification do not more asymmetrical modification in the sequence, where the modification do not not not a device the modification decreasing (1); involves selecting a sequence with activity and introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (1) and instruction for its use; and a device that can be dispense or administer a composition comprising (1). (1) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. (M1) (context) sequence is suffering from a disorder characterised by elevated or cherwise unwanted expression of apoB-100, elevated or cherwise unwanted levels of cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from the HDL/LDL cholesterol imbalance, hypercholesterolaemia, statin-resistant characteriae cholesterolaemia, coronary artery disease (CAD), coronary heart disease (CHD) and atherosclerosis. (1) is administered to a subject to inhibit hepatic glucose production or for treating the diseases as mentioned above, cancer (e.g. breast, colon or clump acheen a hepatitis C viral disease (e.g., AIDS). This sequence represents a hepatitis C viral (120) antisense oligonucleotide that can control HCV gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense sequence and antisense sequence which has specific modifications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention describes a RNA interference (iRNA) agent (I) comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; SEQ ID NO 6756; 378pp; English.
                                                                  14-APR-2003; 2003US-0462894P.
17-APR-2003; 2003US-0463772P.
25-APR-2003; 2003US-046565F.
25-APR-2003; 2003US-0465602P.
09-MAY-2003; 2003US-0469612P.
                                                     2003US-0455050P.
                                                                                                                                                                                       08-AUG-2003; 2003US-0493986P.
11-AUG-2003; 2003US-0494597P.
                                                                                                                                                                                                                                   2003US-0506341P.
2003US-0510246P.
  2003US-0454265P
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                                                                                                                                                                                                                                                                                                                                                                                                       Bumcrot D;
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                      13-MAR-2003; 2
13-MAR-2003; 2
14-APR-2003; 2
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 0.7%; Score 19; DB 1; Length 19;
00.0%; Pred. No. 7.7e+02;
ve 0; Mismatches 0; Indels
0./c,
100.0%; Pre
                          19; Conservative
Query Match
Best Local Similarity
                          Matches
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ò 셤 RESULT 826

ADR82261 standard; DNA; 19 ADR82261; ADR82261 SXXX

BP

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cytostatic; anticonvulsant; notropic; muscula; anti-HIV; Mark interference; IRN; antisense technology; lipid metabolism; cholesterol imbalance; dyslipidaemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; disease; cancer; bung cancer; hung cancer; hung cancer; neurological disease; Huntington disease; spinocerebellar ataxia; viral disease; ALDS; hepatitis C virus; HCV; ss.
                                        cardiant; vasotropic; antiarteriosclerotic; antidiabetic;
                   Hepatitis C virus (HCV) oligonucleotide segid 6760.
                                                                                                                                                                                                                                   12-WAR-2003; 2003US-0454265P.
13-WAR-2003; 2003US-0454962P.
13-WAR-2003; 2003US-0454962P.
17-APR-2003; 2003US-0462894P.
17-APR-2003; 2003US-046565P.
25-APR-2003; 2003US-046565P.
09-MAY-2003; 2003US-046565P.
11-AUG-2003; 2003US-049459P.
26-SEP-2003; 2003US-050541P.
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2003US-0454265P.
2003US-0454962P.
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                                                                                                                                                                                                                                                                                                                                                             10-OCT-2003; 2003US-0510318P
 (first entry)
                                                                                                                                          Hepatitis C virus.
                                                                                                                                                               WO2004080406-A2.
                                       antilipemic;
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 16-DEC-2004
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(ALNY-) ALNYLAM PHARM

Manoharan M, Bumcrot D;

WPI; 2004-677362/66.

Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense sequence and antisense sequence which has specific modifications.

Example 5; SEQ ID NO 6760; 378pp; English.

come or more asymmetrical 2'-0 alkyl modifications, the sense sequences have one or more asymmetrical 19-0 alkyl modifications, the antisense cone or more asymmetrical phosphorothioate modifications cone or more asymmetrical phosphorothioate modifications and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (1); reducing (M1) apoB-100 cc levels or glucose-6-phosphatase levels in a subject; producing (II); clavels or glucose-6-phosphatase levels in a subject; producing (II); clavels or more asymmetrical modification in the sequence, where introducing one or more asymmetrical modification in the sequence, where crete modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (I) and instruction for its use; and a device that can be dispense or administer a composition comprising (I). (I) is activity; a kit comprising (I) and instruction for its use; and a device that can be dispense or administer a composition comprising (I). (I) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. (M1) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. (C) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. (C) cherwise unwanted expression of apoB-100, elevated or otherwise unwanted correspondent or dispense or type-2 dispense (GD) inhibit hepatic glucose production or for treating glucose-engage or dispense or type-2 invention describes a RNA interference (iRNA) agent (I) comprising a

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                                                                                                                                                                                                                                                                                                                                                                                                                    antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; cytostatic; anticonvulsant; nootropic; muscula; anti-HIV; MAN interference; iRNA; antisense technology; lipid metabolism; cholesterol imbalance; dyslipidaemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; diabetes; cancer; breast cancer; spinocencer; lung cancer; neurological disease; Huntington disease; spinocerebellar ataxia; viral disease; ADDS; hepatitis C virus; HCV; ss.
treating the diseases as mentioned above, cancer (e.g. breast, colon or lung cancer), neurological disease (e.g., Huntington disease or spinocerbellar ataxia) or viral disease (e.g., AIDS). This sequence represents a hepatitis C virus (HCV) antisense oligonucleotide that can be used to control HCV gene expression.
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                                                                                                                        DB 1; Length 19;
7.7e+02;
hes 0; Indels
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                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                        Best Local Similarity 100.0%; Pred. No. 7.7
Matches 19; Conservative 0; Mismatches
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                                                                                                                            0.7%; Score 19;
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2003US-0493986P.
2003US-0494597P.
2003US-0506341P.
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2003US-0463772P.
2003US-0465665P.
2003US-0465802P.
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2003US-0454265P.
2003US-0454962P.
2003US-0455050P.
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                                                                                                                                                                                                                                                                                                                                                             16-DEC-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus.
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08-AUG-2003;
11-AUG-2003;
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25-APR-2003;
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                                                                                                                            Query Match
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The invention describes a RNA interference (IRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequences have one or more asymmetrical 2'-0 alkyl modifications, the antisense cone or more asymmetrical phosphorothioate modifications and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (I); reducing (MI) apoB-100 crevels or glucose-6-phosphatase levels in a subject; producing (I); involves selecting a sequence with activity and introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (I) and instruction for its use; and a device that can be disponse or administer a composition comprising (I). is useful for reducing apoB-100 levels or glucose-6-phosphatase levels.

Cuseful for reducing apoB-100 levels or glucose-6-phosphatase levels.

The subject is suffering from a disorder characterised by elevated or otherwise unwanted expression of apoB-100, elevated or otherwise unwanted classes of cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from the HDL/LDL cholesterol imbalance, disorder is chosen from the HDL/LDL cholesterol imbalance, disorder is chosen procession of apoB-100, elevated to a subject to inhibit hepstic glucose production or for treating glucose-metabolism. The disease (CHD) and atherosclerosis. (I) is administered to a subject to inhibit hepstic glucose production or for treating glucose-metabolism. related disorder e.g. diabetes or type-2 diabetes. (I) is useful for treating the diseases as mentioned above, cancer (e.g. breast, colon or treating the diseases as mentioned above, cancer (e.g. breast, colon or treating the diseases as mentioned above, cancer (e.g. breast, colon or ungenesses) a hepatitis C viral (HCV) antisense oligonucleotide that can ö antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; cytostatic; anticonvulsant; nootropic; muscula; anti-HIV; RNA interference; iRNA; antisense technology; liptid metabolism; cholesterol imbalance; dyslipidaemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; diabetes; cancer; breast cancer; colon cancer; lung cancer; neurological disease; Huntington disease; spinocerebellar ataxia; viral disease; ADDS; hepatitis C virus; HCV; ss. Gaps ö 0.7%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 7.7e+02; 0; Indels Hepatitis C virus (HCV) oligonucleotide segid 6755. Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other; 100.0%; Prec. ... 2709 AAAAAAAAAAAAAAA 2727 ВР 07-MAR-2003; 2003US-0452682P. 12-MAR-2003; 2003US-0454265P. 13-MAR-2003; 2003US-0454962P. 13-MAR-2003; 2003US-04550509P. 14-APR-2003; 2003US-0462894P. 38-MAR-2004; 2004WO-US007070 ADR82256 standard; DNA; 19 (first entry) 19; Conservative Best Local Similarity Hepatitis C virus. WO2004080406-A2 16-DEC-2004 23-SEP-2004. ADR82256; Query Match RESULT 828 ADR82256/c Matches ð

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PR 17-ARP-2003; 2003US-046565EP.
PR 25-ARP-2003; 2003US-046565EP.
PR 62-ARP-2003; 2003US-0465612P.
PR 62-ARP-2003; 2003US-0465612P.
PR 62-ARP-2003; 2003US-0465612P.
PR 62-ARP-2003; 2003US-046591P.
PR 62-ARP-2003; 2003US-046541P.
PR 62-ARP-2003; 2003US-051034EP.
PR 11-AUG-2003; 2003US-051031EP.
PR 10-OTT-2003; 2003US-05103IEP.
PR
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Query Match 0.7%; Score 19; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 7.7e+02; Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps
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## 2709 AAAAAAAAAAAAAAAA 2727 ||||||||||||||| 19 AAAAAAAAAAAAAAA 1

8 6

antilipemic; cardiant; vascropic; antiarteriosclerotic; antidiabetic; cytostatic; anticonvulsant; nootropic; muscula; anti-HIV; RNA interference; intonvulsant; nootropic; muscula; anti-HIV; RNA interference; intonvulsant; nootropic; muscula; anti-HIV; cholesterolimbalance; dyslipidaemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepetic glucose production; glucose-metabolism-related disoxder; diabetes; cancer; breast cancer; spinoce-metabolism-related disoxder; diabetes; neurological disease; Huntingron disease; spinocerebellar ataxia; viral disease; AIDS; hepatitis C virus; HCV; ss. WO2004080406-A2.

23-SEP-2004; 2004WO-US007070.

07-MAR-2003; 2003US-045265P.

12-MAR-2003; 2003US-045365P.

13-MAR-2003; 2003US-045305P.

11-MAR-2003; 2003US-045305P.

11-ARR-2003; 2003US-045305P.

25-ARR-2003; 2003US-045305P.

25-ARR-2003; 2003US-045305P.

26-SEP-2003; 2003US-045305P.

26-SEP-2003; 2003US-045305P.

26-SEP-2003; 2003US-045305P.

26-SEP-2003; 2003US-0510318P.

07-NOV-2003; 2003US-0510318P.

Manoharan M, Bumcrot D;

WPI; 2004-677362/66.

Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense sequence and antisense sequence which has specific modifications.

Example 5; SEQ ID NO 6758; 378pp; English

The invention describes a RNA interference (iRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequences have one or more asymmetrical 2'-0 alkyl modifications, the antisense can demonstrated to a sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (I); reducing (M1) apoB-100 levels or glucose-6-phosphatase levels in a subject; producing (I); reducing (I); involves selecting a sequence with activity and introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (I) and instruction for its use; and a device that can be dispense or administer a composition comprising (I). (I) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels.

The subject is suffering from a disorder characterised by elevated or cherwise unwanted expression of apoB-100, elevated or otherwise unwanted expression of apoB-100, elevated or otherwise unwanted disorder is chosen from the HDL/DD. cholasterol imbalance, dyslipidaemias, hypercholestorolasmia, statin-resistant for cholesterolaemia, coronary artery disease (CAD), coronary heart of sisease (CHD) and atherosclerosis. (I) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism.

The subject diseases as mentioned above, cancer (e.g. breast, colon or lung cancer), neurological disease (e.g., Hutrington disease or capmocration or interesting succession or spinocerbellar ataxia) or viral disease (e.g., AIDS). This sequence contractive coronary artery disease (e.g., AIDS). This sequence

ngs.res

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ADT85248;
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ADT85248/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New interference RNA agent comprising sense sequence and antisense sequence having cholesterol moieties, useful for reducing apoB-100 levels
                                                                                                                                                                                                                                  antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; familial combined hyperlipidaemia; acquired hyperlipidaemia; hypercholesterolaemia; statin-resistant hypercholesterolaemia; coronary artery disease; coronary heart disease; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; type-2 diabetes; glitaxzone-resistant diabetes; HCV; hepatitis C virus; antisense inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes an interference RNA (iRNA) agent (I) comprising sense sequence and an antisense sequence, where the sense sequence comprises one or more cholesterol moieties, and the antisense sequence targets a human gene sequence. The following are disclosed: a
                                                            Gaps
                                                            ö
                                                                                                                                                                                                                Hepatitis C virus (HCV) inhibition associated DNA seqid 5285.
                                       Length 19;
                                                          Indels
                    Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                                      0.7%; Score 19; DB 1; Le
100.0%; Pred. No. 7.7e+02;
ative 0; Mismatches 0;
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be used to control HCV gene expression.
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2003US-0462894P.
2003US-0465665P.
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                                                                                                                                                     ADT85243 standard; DNA; 19
                                                                                                                                                                                              (first entry)
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                                                Local Similarity 100.
ses 19; Conservative
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                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus.
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17-APR-2003;
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pharmaceutical composition comprising (1); and a device for administering (1) into a patient. (1) is useful for reducing apoB-100 levels or glucose -6-phosphatase levels in a subject. (1) targets a sequence identical to any one of sequences as given in the specification. (1) comprises a cholesterol moiety is coupled to a sense strand. (2) further comprises a second cholesterol moiety. The second cholesterol moiety is coupled to a sense strand. (1) has 10 or more nucleotides. The duplex region of (1) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted evels of cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from HDL/LDL cholesterol imhalance, dyslipidaemia (e.g., familial combined hyperlipidaemia or acquired hyperlipidaemia, coronary the cholesterolaemia, statin resistant hypercholesterolaemia, coronary at the cholesterolaemia statin resistant hypercholesterolaemia, coronary at the cholesterolaemia statin resistant hypercholesterolaemia, coronary at the cholesterolaemia statin resistant hypercholesterolaemia. ô artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesterolaemia. (I) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism-related disorders e.g., type-2 diabetes or glitaxone-resistant diabetes. (I) has endonuclease or exonuclease resistance. This sequence represents a human hepatitis C virus polynucleotide associated with the inhibition of HCV in human liver cells. antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; familial combined hyperlipidaemia; acquired hyperlipidaemia; hypercholesterolaemia; statin-resistant hypercholesterolaemia; coronary artery disease; coronary heart disease; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; type-2 diabetes; glitaxzone-resistant diabetes; HCV; hepatitis C virus; antisense inhibition; ss. Gape ö Hepatitis C virus (HCV) inhibition associated DNA segid 5290. Length 19; 0; Indels Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other; Score 19; DB 1; L Pred. No. 7.7e+02; 100.0%; Pred. No. 100.0%; Pred. No. Mismatches 2709 AAAAAAAAAAAAAAAAA 2727 19 AAAAAAAAAAAAAAA 1 2003US 0461915P 2003US -0462894P 2003US -0463802P 2003US -0463802P 2003US -0463802P 2003US -0493986P 2003US -0494597P 2003US -0494597P 2003US -0506541P 26-SEP-2003; 2003US-0506341P. 09-OCT-2003; 2003US-0510246P. 10-OCT-2003; 2003US-0510318P. 09-APR-2004; 2004WO-US011255. 2003US-0462097P. 0.7%; ADT85248 standard; DNA; 19 (first entry) Local Similarity 100. nes 19; Conservative Hepatitis C virus. WO2004091515-A2. 10-APR-2003; 2 14-APR-2003; 2 17-APR-2003; 2 25-APR-2003; 2 25-APR-2003; 2 09-MAY-2003; 2 08-AUG-2003; 11-AUG-2003; 09-APR-2003; 13-JAN-2005 28-OCT-2004.

09-APR-2004; 2004WO-US011255

Hepatitis C virus. WO2004091515-A2. 28-OCT-2004.

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The invention describes an interference RNA (iRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequence comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence categors a human gene sequence. The following are disclosed: a pharmaceutical composition comprising (I); and a device for administering complements of sequences as given in the specification. (I) comprises a cholesterol moiety is coupled to a sense strand.

(I) further comprises a second cholesterol moiety. The second cholesterol moiety is coupled to a sense strand.

(I) further comprises a second cholesterol moiety. The second cholesterol moiety is coupled to a sense strand.

(I) further comprises a second cholesterol moiety. The second cholesterol complexity is 19 mucleotides in length. The second cholesterol complexity is 19 mucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted levels of expression of (I) is 19 mucleotides in length. The subject is suffering from a disorder having elevated or or otherwise unwanted cholesterol complexity is combined hyperlipidaemia or acquired hyperlipidaemia, statin-resistant hypercholesterolaemia, statin-resistant hypercholesterolaemia, statin-resistant disease, coronary artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesterolaemia. (I) is administered to a subject columbia hepatic glucose production or for treating glucose-metabolism-calarded and human hepaticis (Virus polymucleotide associated with the inhibition of the human is a human hepaticis.
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                                                                                                                                                                                                                                                                     New interference RNA agent comprising sense sequence and antisense sequence having cholesterol moieties, useful for reducing apoB-100 levels or glucose-6-phosphatase levels.
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                                                                                                                                                          Elbashir S, Harborth J;
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ID ADT85244 standard; DNA; 19 BP.
07-NOV-2003; 2003US-0518453P.
08-MAR-2004; 2004WO-US007070.
05-APR-2004; 2004WO-US010586.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of HCV in human liver cells.
                                                                                                          (ALNY-) ALNYLAM PHARM INC.
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comprises one or more cholesterol moieties, and the sense sequence comprises one or more cholesterol moieties, and the antisense sequence targets a human gene sequence. The following are disclosed: a comprise to pharmaceutical composition comprising (1); and a device for administering pharmaceutical composition comprising (1); and a device for administering (2) into a patient. (1) is useful for reducing apos-100 levels or glucose of phosphatase levels in a subject. (1) targets a sequence identical to any one of sequences as given in the specification. (1) comprises a coholesterol moiety is coupled to a sense strand. (2) further comprises a second cholesterol moiety. The subject is moiety is coupled to a sense strand. (1) has 21 or more nucleotides. The ouplex region of (1) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted evels of expression of apo-8-100, elevated or otherwise unwanted levels of cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from HDL/LDL cholesterol imbalance, dyslipidaemia (e.g., familial combined hyperlipidaemia or acquired hyperlipidaemia, statin-resistant hypercholesterolaemia, cononary
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10-APR-2003; 2003US-0461915P.

17-APR-2003; 2003US-0462894P.

25-APR-2003; 2003US-0465665P.

25-APR-2003; 2003US-0465665P.

09-MAY-2003; 2003US-0465802P.

09-MAY-2003; 2003US-0499396P.

11-AUG-2003; 2003US-0499396P.

26-SEP-2003; 2003US-049939P.
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10-OCT-2003; 2003US-0510318P.
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                                                                                                                                                                                                                                                                                                                                                                  interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; demilial combined hyperlipidaemia; acquired hyperlipidaemia; coronary artery disease; coronary heart disease; atherosclerolaemia; coronary artery disease; coronary heart disease; atherosclerosis; type-2 diabetes; glicaxzone-resistant disease; atherosclerosis; type-2 diabetes; glicaxzone-resistant diabetes; HCV; hepatitis C virus; antisense inhibition; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Manoharan M, Elbashir S,
                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ALNY-) ALNYLAM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-766693/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus.
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25-APR-2003;
09-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-SEP-2003;
09-OCT-2003;
10-OCT-2003;
                                                                                                                                                                                                                                                                                                                                                       antilipemic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-NOV-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-AUG-2003;
                                                                                                                                                                                                              13-JAN-2005
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                                                                                                                                         ADT85246;
       RESULT 833
                                               ADT85246,
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hypercholesterolaemia, statin-resistant hypercholesterolaemia, coronary artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesterolaemia. (i) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism-related disorders e.g., type-2 diabetes or glitaxzone-resistant diabetes. (I) has endonuclasse or exonuclasse resistant diabetes a human hepatitis C virus polynuclassic resistant diabetes. Human hepatitis C virus polynuclassic associated with the inhibition of HCV in human liver cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; familial combined hyperlipidaemia; acquired hyperlipidaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypercholesterolaemia; statin-resistant hypercholesterolaemia; coronary artery disease; coronary heart disease; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; type-2 diabetes; glitaxzone-resistant diabetes; HCV; hepatics antisense inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus (HCV) inhibition associated DNA seqid 5287.
                                                                                                                                                                                                                                                                                                                                                       Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-2003; 2003US-0462097P.
10-APR-2003; 2003US-0461915P.
17-APR-2003; 2003US-0461915P.
17-APR-2003; 2003US-0465665P.
25-APR-2003; 2003US-0465665P.
09-MAY-2003; 2003US-0465665P.
09-MAY-2003; 2003US-0493986P.
11-AUG-2003; 2003US-0493986P.
11-AUG-2003; 2003US-0493986P.
10-OCT-2003; 2003US-050341P.
09-OCT-2003; 2003US-0510318P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19 AAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADT85245 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Elbashir S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JAN-2005 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                           19;
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                                 The invention describes an interference RNA (IRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequence comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence cc targets a human gene sequence. The following are disclosed:

targets a human gene sequence. The following are disclosed:

targets a human gene sequence. The following are disclosed:

comprises one or more momental for reducing apoB-100 levels or glucose comparate levels in a subject. (I) targets a sequence identical to any one of sequences as given in the specification. (I) comprises a cholesterol moiety is coupled to a sense strand.

Complex region of (I) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted complex region of (I) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted evels of cholesterol, and/or disregulation of lipid metabolism. The disorder is combined hyperlipidaemia or acquired hypercholesterolaemia, (C) hypercholesterolaemia, statin-resistant hypercholesterolaemia, statin-resistant theoreticosis, preferably creating resistant hypercholesterolaemia, (I) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism.

Conditions of sequence are sestance. This sequence represents a human hepatitis C virus polynuclease resistance. This sequence represents con thuman hepatitis C virus polynuclease resistance. This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; familial combined hyperlipidaemia; acquired hyperlipidaemia; hypercholesterolaemia; statin-resistant hypercholesterolaemia; statin-resistant hypercholesterolaemia; hypercholesterolaemia; cononary heart disease; atherosclerosis; hepatic glucose production; glucose-metabolism-related disorder; type-2 diabetes; glitaxzone-resistant diabetes; HCV; hepatitis C virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus (HCV) inhibition associated DNA segid 5289.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match

Best Local Similarity 100.0%; Pred. No. 7.7

Matches 19; Conservative 0; Mismatches
              Example 4; SEQ ID NO 5287; 324pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 AAAAAAAAAAAAAAA 1
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10-APR-2003; 2003US-0461915P.
14-APR-2003; 2003US-0462894P.
17-APR-2003; 2003US-046565P.
25-APR-2003; 2003US-046565P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-APR-2004; 2004WO-US011255.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JAN-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADT85247;
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comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence targets a human gene sequence. The following are disclosed: a pharmaceutical composition comprising (1); and a device for administering (1) into a patient. (1) is useful for reducing apoB-100 levels or glucose e-phosphatase levels in a subject. (1) targets a sequence identical to any one of sequences as given in the specification. (1) comprises a cholesterol moiety. The coupled to a sense strand. (1) further comprises a second cholesterol moiety. The second cholesterol moiety is coupled to a sense strand. (1) has coupled to a sense strand. (1) has considered a sense strand. (2) cuplex region of (1) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted exposession of apo-B-100, elevated or otherwise unwanted levels of cholesterol, and/or disregulation of lipid metabolism. The disorder is combined hyperlipidaemia or acquired hyperlipidaemia (e.g., familial combined hyperlipidaemia or acquired hyperlipidaemia).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New interference RNA agent comprising sense sequence and antisense sequence having cholesterol moieties, useful for reducing apob-100 levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to inhibit hepatic jucose production or for treating glucose-metabolism-related disorders e.g., type-2 diabetes or glitaxzone-resistant diabetes (I) has endonuclease or exonuclease resistance. This sequence represents a human hepatitis C virus polynucleotide associated with the inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypercholesterolaemia, statin-resistant hypercholesterolaemia, coronary artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesterolaemia. (I) is administered to a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes an interference RNA (iRNA) agent (I) comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO 5289; 324pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Manoharan M, Elbashir S, Harborth J;
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                                                                         26-SEP-2003; 2003US-0506341P.
09-0CT-2003; 2003US-0510246P.
10-0CT-2003; 2003US-0510318P.
07-NOV-2003; 2003US-0516453P.
08-MAR-2004; 2004WO-US007070.
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2003US-0493986P
                                           11-AUG-2003; 2003US-0494597P
                                                                                                                                                                                                                                                                                    35-APR-2004; 2004WO-US010586
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                                                                                                                                                                                                                                                                                                                                                                           (ALNY-) ALNYLAM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-766693/75.
08-AUG-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-SEP-2005
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The invention relates to a novel chemically synthesized double stranded short interfering nucleic acid molecule that directs cleavage of a signal transducer and activator of transcription 3 (57473) RNA by RNA contransducer and activator of transcription 3 (57473) RNA by RNA contransducer and activator of turther includes a composition comprising the short interfering nucleic acid has cytostatic, antiporiatic, dermatological, antinflammatory, and gastrointestinal-Gen. activities. The short interfering nucleic acid or its composition is useful for treating, inhibiting, or reducing cancer, proliferative, and/or preventing, inhibiting, or reducing cancer, proliferative, and/or such as psoriasis, eczema, disorders, or conditions in a subject or organism, such as psoriasis, eczema, dermatilis, Crohn's disease, and inflammatory bowel disease, and for any other disease, trait, or condition that is celated to or will respond to the levels of STT31 in a cell or tissue, alone or in combination with other treatments or therapies. This oligo sequence represents a STAT-3 siRNA strand of the invention.
The invention relates to nucleosides/oligonucleotides (I) modified at 2'

CC position by linker-bound support including nucleosides containing

CC position by linker-bound support including nucleosides containing

CC therapeutics and as research reagents and kits. The method is useful in diagnostics,

the preparation of antisense oligonucleotides that can be directed

CC the preparation of antisense oligonucleotides that can be directed

CC dagainst a target messenger RNA sequence or alternatively against a target

CC omplementary. They are useful for treating organisms having a disease

CC chemical and biophysical properties for cellular membrane penetration

C. they are capable of improving cellular lipid bilayers penetrating

CC chemical as wall as resistance to exonuclease and endonuclease

CC degradation in vivo. (I) mitigate potential problems such as very short

CC biological half-lives due to degradation by nucleases, inherent negative

CC charge and hydrophilic natural composition. The bioreversible protecting

CC cligonucleotides of natural composition. The bioreversible protecting

CC charge and hydrophilic natural sequence to the oligonucleotides and are removed

CC nuclean and hydrophile that in the cell cytosol extract by

CC charge and properties the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleoside/oligonucleotides modified at 2' position by linker-bound support including nucleosides containing bioreversible phosphate blocking linkages, useful as diagnostic, therapeutic and research reagents.
                                                                                                          internucleotide linkages between nucleotides 1-2,2-3,3-4,16-17,17-18,18-19 OR 2-(pivaloylthio)ethyl
phophotritester internucleotide linkages between 1-2,2-3,3-4-4-5,15-16,16,17,17-18,18-19 where nucleotides 1-4 and 15-18 are 2'-0-(2-methoxyethyl)thymidines"
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                                                                             /mod_base= OTHER
/note= "2-(pivaloylthio)ethyl phophotriester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 19; DB 1; Length 19;
.00.0%; Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
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                              Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                             Guzaev A;
                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                             modified base
                                                                                                                                                                                                                                                                                                                                                                               Manoharan M,
                                                                                                                                                                                                                                                                               10-JUN-1999;
                                                                                                                                                                                                                JS6919437-B1
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   Synthetic
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Novel chemically synthesized double stranded short interfering nucleic acid molecule that directs cleavage of STAT3 RNA through RNA interference, useful for treating cancer and inflammatory diseases e.g.

Example 3; SEQ ID NO 551; 266pp; English.

psoriasis in subject or organism

SIRN-) SIRNA THERAPEUTICS INC.

Mcswiggen J;

Robin H,

WPI; 2005-604649/62.

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Signal-transducer and activator of transcription-3; RNA interference; gene silencing; cytostatic; antipsoriatic; dermatological; antiinflammatory; gastrointestinal-Gen.; cancer; inflammation; psoriasis; eczema; dermatitis; Crohns disease; inflammatory bowel disease; siRNA; short interfering RNA; ss.
STAT-3 siRNA antisense strand, SEQ ID 551.
                                                                                                                                                                                                                                                                                                      2001US-0292217P.
2001US-0306883P.
2001US-0311865P.
2002US-0358580P.
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2002US-0386782P.
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2003US-0440129P.
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2003US-00427160.
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2004US-0543480P.
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                                                                                                                                                                                                                                                                     15-DEC-2004; 2004US-00014373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2004US-00826966
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11-MAR-2002;
17-MAY-2002;
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24-NOV-2003;
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06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-2003;
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                                                                                                                                                      Synthetic
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Mcswiggen JA;
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                                                                                                                                                                            Query Match
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Matches
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                                                                                                                                                                        Signal-transducer and activator of transcription-3; RNA interference, gene silencing; cytoetactic; antipsoriatic; dermatological; antiinflammatory; gastrointestinal-den; cancer; inflammation; psoriasis; eczema; dermatitis; Crohns disease; inflammatory bowel disease; siRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel chemically synthesized double stranded short interfering nucleic acid molecule that directs cleavage of STAT3 RNA through RNA interference, useful for treating cancer and inflammatory diseases e.g. psoriasis in subject or organism.
                                  Gaps
                                  ..
                Score 19; DB 1; Length 19;
Pred. No. 7.7e+02;
                                 0; Indels
BP; 19 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                          STAT-3 siRNA target/sense strand, SEQ ID 274.
                     100.0%; Pred. ...
                                                2709 AAAAAAAAAAAAAAA 2727
                                                          (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                         2002US-00151116.
2002WG-018015876.
2002US-0386782P.
2002US-00201394.
2002US-0406784P.
2002US-0406378P.
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2003US-00693059.
2003US-00720448.
2003US-00727780.
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2003WO-US005346.
2003US-00427160.
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2002US-0363124P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-MAY-2004; 2004WO-US016390
                 0.78;
                                                                                                        AEC90676 standard; RNA; 19
                                                                                                                                        17-NOV-2005 (first entry)
                                                                                                                                                                                                          short interfering RNA; ss.
                Query Match
Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J;
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17-MAY-2002;
06-JUN-2002;
22-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                   15-JAN-2003;
20-FEB-2003;
20-FEB-2003;
30-APR-2003;
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10-FEB-2004;
13-FEB-2004;
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23-OCT-2003;
24-NOV-2003;
03-DEC-2003;
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11-MAR-2002;
Sequence 19
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                                                                                                                                                                                                                          Synthetic.
                                                                                                                        AEC90676;
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                                                                                        RESULT 838
                                                                                                 AEC90676/
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                                                                                                             short interfering nucleic acid molecule that directs cleavage of a signal transducer and activator of transcription 3 (STAT3) RNA by RNA short interference. The invention further includes a composition comprising the short interfering nucleic acid has cytostatic, antipsoriatic, dermatological, antipital amentory, and gastrointestinal-Gen activities. The short interfering nucleic acid has cytostatic, antipsoriatic, dermatological, antipital amentory, and gastrointestinal-Gen activities. The short interfering nucleic acid or its composition is useful for treating, preventing, inhibiting, or reducing cancer, proliferative, and/or inflammatory disease, dermatitis, Crohn's disease, and inflammatory bowel disease, and for any other disease, trait, or condition that is related to or will respond to the levels of STAT3 in a cell or tissue, sequence represents a STAT-3 siRNA strand of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New chemically modified double stranded short interfering nucleic acid (siRNA) molecule that directs cleavage of a NOGO receptor (NOGOr) RNA via
                                                                                The invention relates to a novel chemically synthesized double stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA interference, gene silencing, short interfering RNA, siRNA, nervous system injury, spinal cord injury, neuroprotective, vulnerary, cerebrovascular ischemia; cerebroprotective, multiple sclerosis; muscular dystrophy, muscular-gen., neuropathy, motor neurone disease, CNS-gen., ataxia; Parkinsons disease; antiparkinsonian; Huntingtons chorea; anticonvulsant, nootropic, dementia; Creutzfeldt Jakob disease; Alzheimers disease; NOGO receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human NOGO receptor target sequence/siRNA sense strand, SEQ:551.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 0 T; 19 U; 0 Other;
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Example 3; SEQ ID NO 274; 266pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-FEB-2001; 2001US-00780533.
05-ARR-2001; 2001US-00827395.
20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0353124P.
03-ARR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-036782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEF76001 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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ses 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2006-190836/20.
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The invention relates to chemically synthesized short interfering nucleic acids (sinAs) which downregulate expression of a NOGO receptor gene by RNA interference. The sinAs may or may not comparise ribonucleotides, can contain deoxyribonucleotides, can be chemically modified and may be double or shagle stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense regions, or alternatively are assembled from a sense oligonucleotide and may be compositions comprising an sinA targeted from NOGO receptor minAn. It interfering RNA (shRNA). The invention also relates to pharmaceutical compositions comprising an sinA targeted to a NOGO receptor gene, sinAs targeted to a NOGO receptor gene, sinAs targeted to a NOGO receptor gene, sinAs targeted to a NOGO receptor gene itself, and expression vectors and host cells comprising an sinA of the invention. In particular, the invention discloses siRNAs (AEF75903-AEF76100 and AEF76101-AEF76110. targeted to the human NOGO-A (KIAA0886) gene of DDBU accession number BO011787, and siRNAs (AEF75903-AEF76100 and AEF76101-AEF76101) targeted to the human NOGO-A (KIAA0886) gene of DDBU accession number AB020693. The siNAs are used to modulate expression of NOGO receptor or NOGO genes in cells, tissue explants or organisms (e.g., by carriety of neurodegenerative conditions such as central nervous system (CNS) injury (e.g., spinal cord injury or stroke), multiple sclerosis (AES), active organisms (e.g., spinal cord injury or stroke), multiple sclerosis (AES), active are also be used in drug screening, diagnosis, therespounds are also be used in drug screening, diagnosis, therespounds are also be used in drug screening, diagnosis, therespector are also and especially Alzheimer's condition and condition, genetic emplayed and and apprinted and especially and organisms of the parametery of the sinAtive organism or and validation, genetic emplayed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nervous system injury; spinal cord injury; neuroprotective; vulnerary; cerebrovascular ischemia; cerebroprotective; multiple sclerosis; muscular dystrophy; muscular-gen.; neuropathy; motor neurone disease; CNS-gen.; ataxia; Parkinsons disease; antiparkinsonian;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of a human NOGO receptor-targeted double-stranded siRNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         which is identical to the NOGO receptor transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA interference; gene silencing; short interfering RNA; siRNA;
RNA interference (RNAi), useful for modulating gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Huntingtons chorea, anticonvulsant, nootropic, dementia,
Creutzfeldt Jakob disease, Alzheimers disease, NGGO receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 19; DB 1; Length 19; 94.7%; Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NOGO receptor siRNA antisense strand, SEQ:650.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 18 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
                                                  Disclosure; SEQ ID NO 551; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2708 TAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reticulon 4 receptor; RTN4R; ss
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Best Local Similarity 94.77
Matches 18; Conservative
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The invention relates to chemically synthesized short interfering nucleic acids (sinNa) which downregulate expression of a NOGO receptor gene by RNA interference. The siNAs may or may not comprise ribonucleotides, can contain deoxyribonuclectides, can be chemically modified and may be double or single stranded. They further comprise sense and antisense cligonuclectide. Specifically, the sinAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The invention also relates to pharmaceutical compositions comprisitions comprisitions comprisitions and short clutcher discloses siNAs targeted to a NOGO receptor gene, siNAs targeted to a NOGO receptor gene of GenBank accession number BC011787, and siRNAs (ABF75903-ABF76100 and ABF76101-ABF76112) targeted to the human NOGO receptor gene of GenBank accession number BC011787, and siRNAs (ABF75903-ABF76100 and ABF76101-ABF76112) targeted to the human NOGO receptor or NOGO genes in cells, tissue explants or organisms (e.g., by creeptor or NOGO genes in cells, tissue explants or organisms (e.g., by creeptor or NOGO genes in cells, tissue explants of or the treatment of a variety of neurodegenerative conditions such as central nervous system (CNS) injury (e.g., spinal cord injury or stroke), multiple sclerosis (MS), muscular dystrophy, chemotherapy-induced neuropathy amyotrophic disease, dementia, Creuzfeldt-Jakob disease and especially Alaheimer's disease. The sinAs may also be used in drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, therapeutic target identification and validation, genetic engineering, therapeutic target identification and gene mapping (e.g., of therapeutic parget identification and gene mapping (e.g., of therapeutic target identification and sendence and especially Alaheimer's therapeutic target identification and sendence and engi
                                                                                                                                                                                                                                                                                                              chemically modified double stranded short interfering nucleic acid NAA) molecule that directs cleavage of a NOGO receptor (NOGOr) RNA via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transgenic plants; altered petal colour; polymerase chain reaction; ss.
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                                                                                                                                                                                                                                                                                                                                         (siRNA) molecule that directs cleavage of a NOGO receptor (NOGC RNA interference (RNAi), useful for modulating gene expression.
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Pred. No. 7.7e+02;
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100.0%; Pred. No. ...
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              09-FEB-2001; 2001US-00780533.
05-APR-2001; 2001US-00827395.
20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
03-APR-2002; 2002US-0360124P.
06-JUN-2002; 2002US-036782P.
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(first entry)
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Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                               (MCSW/) MCSWIGGEN J A.
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27-APR-1994
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Analysis of cDNA and gene expression - by amplification of mRNA followed
and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double-stranded conns. by using an aggregate of mRNAs and a plural type clabelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction erayme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                            Length 20;
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                                                                                                                     Sequence 20 BP; 1 A; 0 C; 2 G; 17 T; 0 U; 0 Other
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Pred. No. 7.9e+02;
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100.0%; Pred. No. '...
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ID AAQ75576 standard; DNA; 20 BP.
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Les 19; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of a PCR primer which was used in polymerase chain reactions for the amplification of cloned cytochrome P450 sequences. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                Nucleic acid isolate encoding flavonoid-3'-hydroxylase - is used create transgenic plants with altered petal colour.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                       Tanaka Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 aggregate; restriction enzyme; ss
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AAQ75575 standard; DNA; 20 BP.
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93AU-00006698.
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Best Local Similarity
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07-JAN-1993;
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                                                    WO9320206-A1
                                                                                         14-OCT-1993
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                  Synthetic.
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Analyzing target nucleic acid sequences, useful for population genetics, drug development and diagnosing cancer, comprises hybridizing triple forming oligonucleotide and probe to target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "Other= Hypoxanthine or Inosine"
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/mod_base= OTHER
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03-NOV-1999; 99US-0163416P.
21-DEC-1999; 99US-0171348P.
07-JUL-2000; 2000US-0216579P.
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                                                                                                                                                                                                                                                                                                     Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reverse phase triplex forming oligonucleotide, RP-TFO;
protected nucleic acid sequence; PNAS; single nucleotide polymorphism;
SNP; short tandem repeat; cancer; Factor V Leiden SNP; ss.
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                                                                               Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                       Reverse transcription primer used in cDNA analysis technique.
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100.0%; Pred. No. 7.9e+02;
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                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 5; 11pp; Japanese.
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base= OTHER
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/mod base= OTHER
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(first entry)
                               04-AUG-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 19; Conservative
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modified_base
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                                                                                                                                                                                                     16-APR-1993;
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07-SEP-2001
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                                                                                                                        Synthetic.
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     AAQ75576;
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Page 415

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Key

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polymorphisms (SNP) identification probe and determining the SNP score. The methods can be used for analysing target nucleic acid sequences, especially genomic DNA sequences, to determine if they contain SNPs or short tandem repeats (STRs). The methods can be used to detect SNPs for use in population genetics, drug development, forensics, cancer, genetic disease research, genomic analysis, diagnostics and therapeutics in humans, plants and animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analyzing target nucleic acid sequences, useful for population genetics, drug development and diagnosing cancer, comprises hybridizing triple forming oligonucleotide and probe to target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     reverse phase triplex forming oligonucleotide; RP-TFO; protected nucleic acid sequence; PNAS; single nucleotide polymorphism; SNP; short tandem repeat; cancer; Factor V Leiden SNP; ss.
                                                                                                                                                                                                   Gaps
                                                                                                             Revised record issued on 09-SEP-2004 : Correction to Feature Table
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                                                                                                                                                                    0.7%; Score 19; DB 1; Length 20;
|5.0%; Pred. No. 7.9e+02;
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                                                                                                                                          Sequence 20 BP; 19 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                            8-aminopurine substituted region of an RP-TFO.
                                                                                                                                                                              95.0%; Preα. ων.
+ive 0; Mismatches
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                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                                                                                                                                                          AAS05715 standard; DNA; 20 BP
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03-NOV-1999; 99US-0163416P.
21-DEC-1999; 99US-0171348P.
07-JUL-2000; 200UUS-0216579P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-NOV-2000; 2000WO-US030534
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                                                                                                                                                                                   Local Similarity
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(OSTE/) OSTE C C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
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Matches
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The sequence is a second reverse phase triplex forming oligonucleotide, RP-TFO (3' to the SNP) used to analyse Factor V Leiden SNP using the method of the invention. The invention relates to analysing target nucleic acid sequences comprising restricting isolated DNA, hybridising at least one triplex forming oligonucleotide (TFO), adding a 3' to 5'

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structure, hybridising the captured structure with a single nucleotide polymorphisms (SNP) identification probe and determining the SNP score. The methods can be used for analysing target nucleic acid sequences, especially genomic DNA sequences, to determine if they contain SNPs or short tandem repeats (STRS). The methods can be used to detect SNPs for use in population genetics, drug development, forensics, cancer, genetic disease research, genomic analysis, diagnostics and therapeutics in humans, plants and animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel pharmaceutical composition, which has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmentic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                  Revised record issued on 09-SEP-2004 : Correction to Feature Table Key
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 exonuclease to form a protected nucleic acid sequence (PNAS) tail
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                                                                                                                                                                                                                                                         0.7%; Score 19; DB 1; Length 20; 95.0%; Pred. No. 7.9e+02;
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                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 0 C; 0 G; 19 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                              0; Mismatches
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S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ88266 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                  Conservative
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Tang L,
                                                                                                                                                                                                                                                                           Similarity
19; Conserv
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Best Local {
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preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinfilamentory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO antisense gene therapy. The composition is useful for treating or ftp.wipo.int/pub/published\_pct\_sequences 

Seguence 20 BP; 17 A; 1 C; 1 G; 1 T; 0 U; 0 Other;

ö Gaps . DB 1; Length 20; 7.9e+02; hes 0; Indels 100.0%; Pred. ... 0.7%; Score 19; Local Similarity 100. les 19; Conservative Query Match Best Loca Matches

2707 CTAAAAAAAAAAAAAA 2725 CTAAAAAAAAAAAAAA 20

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ABZ89546 standard; DNA; 20 BP. ABZ89546; RESULT 848 

(first entry) 17-0CT-2003 Human oligonucleotide sequence

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiatethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, Sandrasagra A, Ka , Shahabuddin S; Tang L, Li Y, Nyce JW, I Miller S,

WPI; 2003-229219/22

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4788; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antinfilammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory selected in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO Sequence 20 BP; 18 A; 0 C; 0 G; 2 T; 0 U; 0 Other; at ftp.wipo.int/pub/published\_pct\_sequences Query Match 88888888888888888

Gaps ö 0.7%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; ive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 19; Conservative

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2708 TAAAAAAAAAAAAAA 2726 TAAAAAAAAAAAAA 20 ~

ð 셤 RESULT 849 **AB**Z99050,

ABZ99050 standard; DNA; 20

ABZ99050;

17-OCT-2003 (first entry)

Human PDE4C oligonucleotide sequence.

antiniflammatory steroid; ubiquinone; antiniflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. antisense; lung dysfunction; nasal airway dysfunction; 

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

ä Aguilar Katz E, Pabalan J, ŝ Shahabuddin Sandrasagra A, Li Y, Sar Tang L, Miller S, Š Nyce

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 14292; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 and semonic flanking regions, 5 and 3 intron-exon junctions, or regions within 2.10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or ansal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antialergic, antiaethmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, receptor, producing pronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fip.wipo.int/published\_pct\_sequences

Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö Query Match 0.7%; Score 19; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 19; Conservative 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAAAAA 2727 AAAAAAAAAAAAAAA 19

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RESULT 850

ABZ88618 standard; DNA; 20 BP 

ABZ88618;

17-OCT-2003

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 3860; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lug or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition ftp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 19 A; 1 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; 0; Indels 100.0%; Pred. ... 2709 AAAAAAAAAAAAAAAA 2727 Local Similarity 100. hes 19, Conservative Query Match Matches

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ABZ89678 standard; DNA; 20 RESULT 851 ABZ89678

ABZ89678;

BP.

(first entry) 17-OCT-2003 Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinflammic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Sandrasagra A, Katz E, , Shahabuddin S; Li Y, Sar, Tang L, Nyce JW,

Aguilar

WPI; 2003-229219/22

Miller S,

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid ubiquinone.

Disclosure; SEQ ID NO 4920; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiashmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences antisense gene therapy. The composition is useful for 88888888888888

Seguence 20 BP; 19 A; 0 C; 0 G; 0 T; 0 U; 1 Other;

ö Gaps ; 0 DB 1; Length 20; 1; Indels 0.7%; Score 19; DB 1; I 95.0%; Pred. No. 7.9e+02; 0; Mismatches Conservative Query Match Best Local Similarity Matches 19; Conserv

ò 셤 ABZ87681 standard; DNA; 20 BP ABZ87681; 

17-OCT-2003 (first entry)

Human oligonucleotide sequence

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Ното варіепв.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Nyce JW, Miller S,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 2923; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5 or 3 end genomic flanking regions, 15 or 3 end genomic flanking regions, processes of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published\_pct\_sequences 8X8888888888X8

Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö Length 20; Indels Score 19; DB 1; L Pred. No. 7.9e+02; 100.0%; Preq. ... 0.7%; Best Local Similarity 100. Matches 19, Conservative Query Match

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RESULT 853 ABZ89677

BP, ABZ89677 standard; DNA; 20

ABZ89677;

(first entry) 17-OCT-2003 Human oligonucleotide sequence.

Human, antisense, lung dysfunction, nasal airway dysfunction, antisinflammatory steroid, ubiquinone, antisinflammatory; antiallergic, antisinflammatic, hypotensive; immunosuppressive, cytostatic, gene therapy; antisense gene therapy; respiratory, lung, adenosine sensitivity; adenosine receptor; bronchodilation, bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. 

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, ŝ Sandrasagra A, L, Shahabuddin Li Y, Sar Tang L, Miller Š, Nyce JW,

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(8) antisense to specific gene(8) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

Disclosure; SEQ ID NO 4919; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antialbergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a

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use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject s tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                             at ftp.wipo.int/pub/published_pct_sequences
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Sequence 20 BP; 19 A; 0 C; 0 G; 0 T; 0 U; 1 Other;

Score 19; DB 1; Length 20; Pred. No. 7.9e+02; 0; Mismatches 1; Indels 0; Mismatches 2709 AAAAAAAAAAAAAAAA 2728 Query Match 0.7%; Best Local Similarity 95.0%; 19; Conservative Matches ઠે 셤

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Gaps

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1 AAAAAAAAAAAAAAAA 20

ABD24848; RESULT 854 ABD24848 

ABD24848 standard; DNA; 20 (first entry) 29-JUL-2004

AI092623-derived oligonucleotide SEQ ID 3860.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Š

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WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleocide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 3860; 763pp; English.

comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung This invention describes a novel composition (a) a first active agent,

The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung contains and/or bronchoconstriction are associated with a disease or condition such as pulmonary visoconstriction, contains and present in the application, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary visoconstriction, contains or cancer.

Contains and the content of the anti-sense oligos corresponding to transplantation rejection, pulmonary infections, branchines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system content any unwanted effects due to it ö dysfunction or cancer and can be anti-sense to the corresponding mRNA Gaps ö Sequence 20 BP; 19 A; 1 C; 0 G; 0 T; 0 U; 0 Other; Similarity Query Match Best Local S 

0.7%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; ive 0; Mismatches 0; Indels 19; Conservative Matches

2709 AAAAAAAAAAAAAAAA 2727 20 AAAAAAAAAAAAAAAA ~

ò ద RESULT 855

BP. ABD32081 standard; DNA; 20

ABD32081;

(first entry) 29-JUL-2004

Human PDE4C-derived oligonucleotide SEQ ID 14292.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhintis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. PACKET PA

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Aguilar Katz E, Pabalan J, Sandrasagra A, Li Y, Sar , Tang L, Nyce JW, I

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WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to

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This invention describes a novel composition (A) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and considered from a gene encoding or regulating coligonucleotides are derived from a gene encoding or regulating cypression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that composition of the kit. The composition of the invention as antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production of trainflammation, allergies oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung creduce the amount of target polypeptide present in the lungs. The disease or condition such as pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory hypertension, allergies, asthma, impeded respiration, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary of the anti-sense oligos corresponding to thymitammation, pulmonary infections, bronchits or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine tont ent to it tesus environment and thereby, to it prevent any unwanted effects due to it
                                                                                                                                                        novel composition (a) a first active agent,
nucleic acids associated with lung airway or lung dysfunction, and
                                                                                         Claim 15; SEQ ID NO 14292; 763pp; English.
                                                                                                                                                              describes a
                                 bronchodilating agent.
   ##X&XBBBBBBBBBBBBBBBBBBBBBBBBBBBB
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Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ; 0 0.7%; Score 19; DB 1; Length 20; 00.0%; Pred. No. 7.9e+02; 0; Indels 0; Mismatches 100.0%; Best Local Similarity 100.0 Matches 19; Conservative Query Match

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ઠ 셤 856

ABD25776 standard; DNA; 20 BP. ABD25776;

(first entry) AI085559 DNA fragment. 29-JUL-2004 

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory districtses syndrome; allergic rhinitis; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; bronchitis; pulmonary transplantation rejection; ds.

Homo sapiens

40200285309-A2.

23-APR-2002; 2002WO-US013143

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating depression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiaflammatory, antiaflammatory, antiaflammatory, antiaflammatory, antiaflammatory, antiaflammatory, antiaflammatory, is a composition comprises oligo and is administered to reduce the production or vaailability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cumposition and/or bronchoconstriction and associated with a disease or condition such as pulmonary disease, pulmonary disease, pulmonary disease, pulmonary disease, pulmonary disease, pulmonary of the range of the anti-sense oligon corresponding to the range breaktion rejection, pulmonary infections, bronchitis or cancer. The rangelantation rejection, pulmonary infections, bronchitis or cancer. The rangelantation rejection, pulmonary infections, bronchitis or cancer. The rangelantation rejection, pulmonary infections bronchitis or cancer. The velocued adenosine content of the arreget RNA serves to prevent any unwanted effects
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                                                                                                                                                                                                                                                 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                       Aguilar D;
                                                                                                                       Pabalan J,
                                                                                                                       Katz E,
                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 4788; 763pp; English
                                                                                                                          i Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                     24-APR-2001; 2001US-0286036P.
                                                                      (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                      bronchodilating agent.
                                                                                                                                                                                                   WPI; 2003-093058/08
                                                                                                                          Li Y,
                                                                                                                                                     Miller S,
                                                                                                                             Nyce JW,
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Sequence 20 BP; 18 A; 0 C; 0 G; 2 T; 0 U; 0 Other;

ö 0.7%; Score 19; DB 1; Length 20; 00.0%; Pred. No. 7.9e+02; 0, Indels 100.0%; Prec. ... 19; Conservative Query Match Best Local Similarity Matches

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Gaps

2708 TAAAAAAAAAAAAAAA 2726 2 TAAAAAAAAAAAAAAA 20 셤 ઠે

RESULT 857

ABD24496

AI652901-derived oligonucleotide SEQ ID 3508. ABD24496 standard; DNA; 20 BP. 29-JUL-2004 (first entry) ABD24496; 

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;

beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Katz E, Pabalan J, Aguilar D; i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Š

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 3508; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprolesses (c) and elivery of the invention as antiallargic, antinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or maligonal disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition and sheases or condition and/or bronchoconstriction and/or lung and sheases or condition such as pulmonary obstruction, and/or bronchoconstriction are associated condition and asserting and asserting and in allergies and/or such as pulmonary or section. transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary

Sequence 20 BP; 17 A; 1 C; 1 G; 1 T; 0 U; 0 Other;

; 0.7%; Score 19; DB 1; Length 20; 00.0%; Pred. No. 7.9e+02; 0; Indels Mismatches 100.08; Pr. Best Local Similarity 100. Matches 19; Conservative Query Match

2707 CTAAAAAAAAAAAAA 2725

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CTAAAAAAAAAAAAA 20

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Gaps

ABD23911/c

ABD23911 standard; DNA; 20 BP.

ABD23911;

(first entry) 29-JUL-2004 Human calmodulin 2-derived oligonucleotide SEQ ID 2923.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; empirysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

'n Aguilar Pabalan J, Katz E, ŝ Sandrasagra A, J, Shahabuddin Li Y, Sar, , Tang L, Nyce JW, I Miller S,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted truncleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 2923; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstruction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, confidence are derived from a gene encoding or regulating reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung advice, in separate containers, (b) the oligonucleotides, (c) instruction and adding carrier and for use of the kit. The composition of the invention has antiallergic, antiniflammatory, antiathmatic, is a classification and expression of the inventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and or cancer. The reduced adenosine content of thoractive pulmonary disease, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc. tissue environment and thereby, to it its prevent any unwanted effects due to it 

Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

ngs.res

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The present invention relates to an oligonuclectide anti-sense to e.g.,
initiation codon, coding region with 2-10 nuclectides of 5'-end and 3'-
end of nucleic acid target comprising gene(s) chosen from e.g.
interleukin (IL)-4 receptor, IL-5 receptor or salts of the
coligonuclectide and optionally surfactant operatively linked to the
oligonuclectide. The method is useful for preventing or treating a
respiratory or lung disease, which involves administering to the airways
of subject an effective amount of an inhibitor. The oligonuclectide is
useful for production of a medicament for the prevention and/or treatment
of a subject an effective amount of an inhibitor. The oligonuclectide is
consen from alrway inflammation, allergy(ses), asthma, impeded
chosen from alrway inflammation, allergy(ses), asthma, impeded
consen from alrway inflammation, allergy(ses), asthma, impeded
(CODP), allergic rhinitis (AR), acute respiratory distress syndrome
(ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway
obstruction. The present sequence represents an oligonuclectide of the
                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease;
airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                           Gaps
                                           ö
  Length 20;
                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Aguilar D, Miller S;
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    DB 1; Le
7.9e+02;
Query Match 0.7%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 7.9 Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated to PDE4C #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; SEQ ID NO 1791; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW, Tang L, Sandrasagra A,
Shahabuddin S, Lu H, Cong H;
                                                                                   2709 AAAAAAAAAAAAAAA 2727
                                                                                                                      19 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                           ADJ60935 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                          06-MAY-2004 (first entry)
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                                                                                                                                                                                                                                                                    ADJ60935;
                                                                                                                                                                                     RESULT 859
                                                                                                                                                                                                          ADJ60935,
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound compositions with and inhibits the expression of Nav1.3. The specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuropathy, arthritic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                           Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                            Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
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100.0%; Pred. No. 7.9e+02;
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2709 AAAAAAAAAAAAAAAA 2727
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                                  ААААААААААААААА 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                          14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-2002; 2002US-0403416P.
                                                                                                                           ADK74647 standard; DNA; 20
                                                                                                                                                                                             (first entry)
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les 19; Conservative
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                                                                                                                                                                                             20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roberds SL;
                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                              ADK74647;
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                                    19
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ID ADK7
XX
AC ADK7
                                                                                                           ADK74647,
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Gaps

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/ Match 0.7%; Score 19; DB 1; Length 20; Local Similarity 100.0%; Pred. No. 7.9e+02; nee 19; Conservative 0; Mismatches 0; Indels

Query Match

Best Loca Matches

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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothhoate oligomicalectide with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound targeted to a nucleic acid molecule encoding abav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                        Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
                                                       Chimeric phosphorothioate oligonucleotide to target Navl.3 #1522.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1748.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 1522; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAA 2727
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                                                                                                                                                                                                                                                                                   14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                      14-AUG-2002; 2002US-0403416P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADK74414 standard; DNA; 20
               20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                        WO2004016754-A2.
                                                                                                                                                                                                                                             26-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                 Roberds SL;
                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK74414;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
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à
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disoater such as childhood seizure disoater, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MoE wings and a deoxy gap. Used during the antisense inhibition of different regions of the human Navi.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; osotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                                                                                                                                                                              New antisense compound targeted to a nucleic acid molecule encoding bay1.3, useful for useful for reating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.7%; Score 19; DB 1; Length 20;
100.0%; Pred. No. 7.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Prea. w.
                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 1748; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAA 2727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 AAAAAAAAAAAAAAA
                                                                                    14-AUG-2003; 2003WO-US025465
                                                                                                                        14-AUG-2002; 2002US-0403416P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM14246 standard; DNA; 20
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Best Local Similarity 100.
                                                                                                                                                         (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                              WPI; 2004-203785/19.
                  WO2004016754-A2.
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                                                   26-FEB-2004.
                                                                                                                                                                                            Roberds SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM14246;
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Gaps

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Location/Qualifiers

Key modified base

Tue Nov

CCRI; CCR3; Eccaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptease; tryptease b; PDB4 A; PDE4 B; PDB4 C; PDB5 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CCP0P; allergic finitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.

88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5;

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiavascular activities, and can ophthalmological, immunomodulatory and cardiavascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                               /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
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100.0%; Pred. No. 7.9e+02;
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                                                                                                         /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                                                          not\overline{e} = "2' - 0 - methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ ID NO 433; 132pp; English.
                 /mod_base= OTHER
                                                                                                                                                                                        /mod base= OTHER
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Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                            25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2002; 2002US-0413549P.
                                                                           ...5
/*tag= a
                                                                                                                                                                       *tag= c
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                                                                                                                                                                                                                                                                                                                                                                                                   (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-305094/28.
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                                                                         modified base
                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                        08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ischemia.
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Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.

Claim 2; SEQ ID NO 1791; 174pp; English.

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Miller

Aguilar D,

Nyce JW, Sandrasagra A, Tang L, Shahabuddin S, Lu H, Cong H;

WPI; 2004-293804/27.

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MILLER S. SHAHABUDDIN AGUILAR D.

(MILL/) (SHAH/) (LUTHH/)

(AGUI/)

CONG H.

23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143. 25-JUL-2003; 2003US-00627930.

US2004049022-A1.

11-MAR-2004.

Homo sapiens

NYCE J W. SANDRASAGRA A.

TANG L.

(SAND/) (TANG/) (NYCE/)

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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL) -4 receptor, interleukin (IL) -5 receptor, CCR1, CCR3, Edeaxin, RANTES, MCP4, CD23, ICAM, WCAM, CCCR3 relates to a method of screening a candidate compound that binds to an enchance of screening a candidate compound that binds to one or more nucleic acid target(8) or expressed product(8), for the prevention and/or treatment of a respiratory or lung disease. The coligonucleotides are useful for reducing or inhibiting expression of a cent or mRNA encoding interleukin 4 receptor, interleukin-5 receptor, CCCR1, CCR3, Ediaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to and/or asthma and/or lawels of adenosine A receptor(8), and/or asthma and/or lawels of adenosine A receptor(8), and/or asthma and/or lawels of adenosine A respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cyptic fibrosis (CP), chronic obstructive pulmanaty or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cyptic fibrosis (CP), chronic obstructive pulmonary disease (COP), chronic obstructive pulmonary disease (COP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 19; DB 1; Length 20;
100.0%; Pred. No. 7.9e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Whiches 19; Conservative
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2709 AAAAAAAAAAAAAAAA 2727

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19 AAAAAAAAAAAAAAA 1

ADO46424 standard; DNA; 20 BP

RESULT 864 AD046424/c Human oligonucleotide #1790

15-JUL-2004 (first entry)

AD046424;

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The invention relates to an isolated human polynucleotide. The composition and methods are useful for screening, diagnosing or treating breast cancer. These may also be used in drug screening or in monitoring disease progression and/or treatment efficacy of breast cancer. The present sequence represents a human small inducible cytokine B14 precursor breast cancer marker cDNA.
Cytostatic; Gene therapy; diagnosis; breast tumor; endocrine disease; gynecology and obstetrics; neoplasm; ss; tumor marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human nucleic acid and polypeptide sequences useful for screening,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pollock S, Levine Z, Akiva P,
O, Ayalon-Soffer M, Walach S;
Keren N, Shklar M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 19; DB 1; Length 20; 00.0%; Pred. No. 7.9e+02;
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Local Similarity 100.0%; Pred. No. (...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 219; 1586pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosing or treating breast cancer.
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ovik A, Sella-Tavor
nwald S, Shemesh R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                     22-OCT-2004; 20040S-0525129; 22-OCT-2004; 20040S-0620656P. 22-OCT-2004; 20040S-0620853P. 22-OCT-2004; 20040S-0620817P. 22-OCT-2004; 20040S-0620917P. 22-OCT-2004; 20040S-0620918P. 22-OCT-2004; 20040S-0620918P. 22-OCT-2004; 20040S-0620974P. 22-OCT-2004; 20040S-062104P. 22-OCT-2004; 20040S-062103P. 17-NOV-2004; 20040S-062131P. 17-NOV-2004; 20040S-0628111P. 17-NOV-2004; 20040S-0628111P. 17-NOV-2004; 20040S-0628111P.
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17-NOV-2004; 2004US-0628134P.
17-NOV-2004; 2004US-0628145P.
17-NOV-2004; 2004US-0628156P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-NOV-2004; 2004US-0628167P.
17-NOV-2004; 2004US-0628178P.
17-NOV-2004; 2004US-0628231P.
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27-JAN-2005; 2005US-00043842.
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                                                                                                                                                                  2005WO-IB000433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diber A, Novik A,
Sameah-Greenwald S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2005-555592/56.
                                                                                          WO2005072050-A2
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                                                      Homo sapiens.
                                                                                                                                                                  27-JAN-2005;
                                                                                                                               11-AUG-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of amplifying an unclonable DNA fragment in vitro for transfection into a eukaryotic cell which comprises amplifying the unclonable DNA fragment by rolling circle amplification (RCA) to produce a tandem series of repeate of the unclonable DNA fragment. The vector is used in therapy, specifically in gene therapy, which may be for unscular dystrophy and for DNA expression studies. The method and kit are useful for amplifying an unclonable DNA fragment in vitro. The present sequence represents an oligo (dT)20 primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amplifying an unclonable DNA fragment in vitro, useful for transfecting into a eukaryotic cell, comprises amplifying the fragment by rolling circle amplification.
                                                                                                                                                                                                                                                                                            Muscular-Gen.; transfection; DNA amplification; gene therapy; growth disorder; musculoskeletal disease; neurological disease; muscular dystrophy; DNA expression; ss; RT-PCR; reverse transcriptase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Pred. No. 7.9e+02;
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100.0%; Pred. No. '...
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Best Local Similarity
Matches 19; Conserv
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gene expression - by amplification of mRNA followed

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GBNESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                  by digestion with restriction enzymes.
                                                                                                          Disclosure, Page 8; 11pp; Japanese
93JP-00112515
                                                                       Analysis of cDNA and
                                                WPI; 1995-018287/03
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16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method (M1) for labeling double stranded nucleic acid for efficient detection of DNA or RNA. The method comprises using a carbodininde derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid. (M1) is useful for labeling double stranded or single stranded nucleic acid or detecting single nucleocide polymorphisms. The present sequence was used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                            Labeling double stranded nucleic acid, involves utilizing carbodiimide derivative for labeling thymine, uracil and guanine, which exists in mismatch region of nucleic acid or unstable region of hydrogen bond of nucleic acid.
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                                                Oligonucleotide ODN3 used to illustrate nucleic acid labeling method
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                                                                         DNA detection; RNA detection; SNP detection; ss
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electrophoresing the digested aggregate of {\tt CDNAs} in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                            0.7%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 8.1e+02; ive 0; Mismatches 0; Indels
                                                                                                    Sequence 21 BP; 1 A; 2 C; 1 G; 17 T; 0 U; 0 Other;
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Best Local Similarity 100.
Matches 19; Conservative
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aggregate; restriction enzyme; ss.
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                                                          01-NOV-1994
                   Synthetic.
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       DB 1; Length 21;
8.1e+02;
hes 0; Indels
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100.0%; Pred. No. 8.1
Live 0; Mismatches
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                                                                                                                                                                                                           aggregate; restriction enzyme; ss.
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AAQ75703 standard; DNA; 21 BP.
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Best Local Similarity 100.
Matches 19; Conservative
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Matches 19; Conservative
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abablled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 8.1e+02;
                                                                                                                                                                                                                                                                             0.7%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 8.1e+02;
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                                                                                                                                                                                                                           Sequence 21 BP; 1 A; 1 C; 2 G; 17 T; 0 U; 0 Other;
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Matches 19, Conservative
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nes 19; Conserv
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19

RESULT 874

Query Match

Matches

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gene expression - by amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                   expression; reverse transcription; primer; cDNA;
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                                                                                                                                                                                                                               Reverse transcription primer used in cDNA analysis technique.
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Pred. No. 8.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Scor.
100.0%; Pred. No. e...
... 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2707 CTAAAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                         aggregate; restriction enzyme; ss
AAQ75707/C

ID

AAQ75707 standard; DNA; 21 BP.

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AAQ75707;
XX
AAQ75707;
XX

O1-AUG-1995 (first entry)
XX
Analysis; gene expression; rever
XX
XX
Analysis; gene expression; rever
XX
Synthetic.
XX
Synthetic.
XX
AN
O1-NOV-1994.
XX

I6-APR-1993; 93JP-00112515.
XX
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I6-APR-1993; 93JP-00112515.
XX
XX
ANALYSIS of CDNA and gene express
PT
CC
CD
Analysis of cDNA and gene express
PT
Analysis of cDNA and gene express
PT
CC
CA double-stranded cDNAs by using a
CC
Transcription primer; (b) digest
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Transcription primer; (c) digest
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Transcription primer; (
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(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
    WPI; 1995-018287/03.
                                            RESULT 876
                                 Matches
                                             AAQ75709,
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ö A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-CD5798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily 당당 gene expression - by amplification of mRNA followed - by amplification of mRNA followed A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESGA files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily Gaps ö Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; 88. Reverse transcription primer used in cDNA analysis technique. Length 21; 0; Indels Sequence 21 BP; 2 A; 1 C; 1 G; 17 T; 0 U; 0 Other; Score 19; DB 1; Le Pred. No. 8.1e+02; 0; Mismatches 0; (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP. by digestion with restriction enzymes. alysis of cDNA and gene expression - digestion with restriction enzymes. Disclosure; Page 7; 11pp; Japanese. Disclosure; Page 7; 11pp; Japanese. 0.7%; Scur 100.0%; Pred 2707 CTAAAAAAAAAAAAAA 2725 19 CTAAAAAAAAAAAAAA 1 BP 93JP-00112515. 93JP-00112515 3709/c AAQ75709 standard; DNA; 21 (first entry) Conservative Analysis of cDNA and WPI; 1995-018287/03. Local Similarity les 19; Conserv JP06303997-A. 16-APR-1993; 16-APR-1993; 04-AUG-1995 Analysis of 01-NOV-1994 AAQ75709; Query Match

Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides (b have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceramic, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, conducting properties and electrical field or through a molecular, hair method is used (i) for analysis of patterns, especially in mucosal, cation is physical, stimulated by an electrical field or through a molecular or additives or supplements, especially minerals, trace elements, organic acids find and activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids find matrix in an acid competed and inacpeted acid in a cell, and can be acids matrix miniaturisation is possible. Abx01281-Abx01344 represent
                                                                                                                                                                                                                                                                                                                                                                               capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for analyzing expression uses capture agent with variable
                                         Gaps
                                       ;
0.7%; Score 19; DB 1; Length 21;
(00.0%; Pred. No. 8.1e+02;
                                     Indels
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                                                                                                                                                                                                                                                                                                                                          Rat DNA microarray capture oligonucleotide #10.
      100.0%; Pred. no.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorting single-stranded nucleic acid, patterns and screening active agents,
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                                                                           2707 CTAAAAAAAAAAAAAAA 2725
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 5; 8pp; German.
                                                                                                                                                                                                                   ADK01290 standard; DNA; 21
                                                                                                                                                                                                                                                                                                    (first entry)
                                       Conservative
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                l Similarity
19; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                             ss; hybridisation;
blood; nerve; germ
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                                                                                                                                                                                                                                                            ADK01290;
Query Match
Best Local {
                                                                                                                                                                                                  ADK01290/c
                  Best Loc
Matches
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capture probes used in the method of the invention.
                                     Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
     SXS
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0.7%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 8.1e+02; tive 0; Mismatches 0; Indels 2708 TAAAAAAAAAAAAAA 2726 Query Match Best Local Similarity 100.0 Matches 19; Conservative 19 8 ò

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Gaps

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RESULT 878 ADK01281/c

ADK01281 standard; DNA; 21 BP. 

ADK01281;

Rat DNA microarray capture oligonucleotide #1. (first entry) 06-MAY-2004

sø; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

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Boekenkamp D, Dieck HT, Hoppe

WPI; 2003-714082/68.

useful for analyzing expression uses capture agent with variable Sorting single-stranded nucleic acid, patterns and screening active agents, and constant regions.

Example; Page 4; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cadents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anthor region. The complement of which is present at least once in each nucleic acid and a 3' variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture of particularly locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by compuring properties are biotinylated and immobilised on a surface capture cliquoucleotides are biotinylated and immobilised on a surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, race elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells.

allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. Sequence 21 BP; 3 A; 0 C; 0 G; 18 T; 0 U; 0 Other; 888888

Gaps . 0 DB 1; Length 21; 0; Indels Score 19; DB 1; Le Pred. No. 8.1e+02; 100.0%; Pred. No. Query Match Best Local Similarity 100.0 Matches 19; Conservative

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2708 TAAAAAAAAAAAAAAA 2726 19 TAAAAAAAAAAAAAA 1 ò g

RESULT 879 ADK01335/

ADK01335 standard; DNA; 21

BP.

ADK01335; 

06-MAY-2004 (first entry)

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #55.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded conclet acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cadents that are (a) immobilised on the surface of a solid matrix and (b) agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprises and non-variable region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow that comprises a sequence of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, interacting properties and especially in the form of a chip. Ints surface conducting properties and especially in the form of a chip. Ints stranded nucleic acids to the surface is (quasi) covalent, supramolecular, stranded nucleic acids to the surface is (quasi) covalent, supramolecular, chair root, blood, nerve or germ cells and (i) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food

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additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of mucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
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Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;

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0
0.7%; Score 19; DB 1; Length 21;
                        0; Indels
            100.0%; Pred. No. 8.1e+02; tive 0; Mismatches 0;
                                                  2709 AAAAAAAAAAAAAAAA 2727
                                                                   19 AAAAAAAAAAAAAAA 1
                         Conservative
            Local Similarity
                        19;
Query Match
             Best Loc
Matches
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ADK01291 standard; DNA; 21 BP 06-MAY-2004 (first entry) ADK01291; RESULT 880 ADK01291/

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #11. 28-FEB-2002; 2002DE-01008794. DE10208794-A1. 04-SEP-2003. Rattus sp. 

Boekenkamp D, Dieck HT, Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794.

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprises as 5-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, interaction with streptavidin. The matrix is of plastic, ceramic, glass, interaction with allowed and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is sparticularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular,

physical, stimulated by an electrical field or through a molecular sieve. The method is used (1) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (1) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.

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Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;

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Gaps

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Gaps
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0
        Score 19; DB 1; Length 21;
Pred. No. 8.1e+02;
                                              0; Indels
Query Match

0.7%; SCOLE 1., __
Best Local Similarity 100.0%; Pred. No. 8.1
Matches 19; Conservative 0; Mismatches
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2708 TAAAAAAAAAAAAAAA 2726 19 TAAAAAAAAAAAAAA 1 ઠે 셤

ВЪ .295/c ADK01295 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01295; RESULT 881 ADK01295, 

Rat DNA microarray capture oligonucleotide #15.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT,

Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5' invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (IMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass,

metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible corpresentation of differences in pools of nucleic acid in a cell, and can detect very small differences in the nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the genes, matrix miniaturisation is possible. ADK01344 represent capture probes used in the method of the invention. 25555555555555555555888

Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 8.1e+02; 0; Indels Query Match
Best Local Similarity 100.0%; Fred. No. 8.10

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ADK01283 standard; DNA; 21 BP 06-MAY-2004 ADK01283; RESULT 882 

(first entry)

Rat DNA microarray capture oligonucleotide #3.

вв; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Dieck HT, Hoppe H; Boekenkamp D,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions. WPI; 2003-714082/68.

Example; Page 4; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture

comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The computions of 10-50, particularly 15-25, T residues. The computed of 10-50 particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly alayer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, compusitoral, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceulucials and/or nutritional compounds, eag food activity of pharmaceulucials and/or nutritional compounds, eag. food activity of pharmaceulucials and/or nutritional compounds, eag food cids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible conditions of differences in pools of nucleic acid from calls. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools not individual genes, matrix miniaturisation is possible. ADX01281-ADX01344 represent 

Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

Gaps ö Length 21; 0; Indels 0.7%; Score 19; DB 1; Le 100.0%; Pred. No. 8.1e+02; iive 0; Mismatches 0; Best Local Similarity 100. Matches 19; Conservative Query Match

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RESULT 883

ADK01339 standard; DNA; 21 ADK01339/ ID ADK0

ВР.

(first entry) 06-MAY-2004

ADK01339;

Rat DNA microarray capture oligonucleotide #59.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1. 

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

ï Hoppe Boekenkamp D, Dieck HT,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides

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that comprises all possible condinations of up to 10 muclecides to allow that comprises all possible combinations of up to 10 muclecides to allow binding of particular sorts of single stranded nucleic acids. The capture binding of particularly locked mucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligomuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. The size of stranded nucleic acids to the surface is (quasi)covalent, supramolecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, additives or supplements, especially minerals, trace elements, organic actids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible captures the method is based on comparison of mucleic acid in a cell, and can detect very small differences in pools of mucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of and locatic acid pools not individual genes, matrix miniaturisation is possible. ADKN1281-ADKN1344 represent

Sequence 21 BP; 0 A; 2 C; 0 G; 19 T; 0 U; 0 Other;

Gaps ö Query Match 0.7%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 8.1e+02; Matches 19; Conservative 0; Mismatches 0; Indels 2709 AAAAAAAAAAAAAAA 2727 19 AAAAAAAAAAAAAA 1 ઠે g

ADK01289 standard; DNA; 21 BP. 06-MAY-2004 (first entry) ADK01289; RESULT 

Rat DNA microarray capture oligonucleotide #9.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

Boekenkamp D, Dieck HT, Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH.

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded

cc reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using capture cagenes that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least nonce in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises are particularly locked nucleic acids. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) molecular filaments and binding of single cranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives. The method provides rapid, inexpensive and reproducible can lows inaging of the complete pattern of all nucleic acids from cells, and can lows inaginal differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, or individual can marrix minaturisation is possible, App(201281-App(01344 represent capture probes used in the method of the invention

Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

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Gaps ö 0.7%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 8.1e+02; iive 0; Mismatches 0; Indels Unery Match Best Local Similarity 100.( Matches 19; Conservative

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ADK01294 standard; DNA; 21 BP. ADK01294; RESULT 885 ADK01294, 

Rat DNA microarray capture oligonucleotide #14.

(first entry)

06-MAY-2004

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Dieck HT, Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Bush A;

Mauck K,

Bartel P,

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ADZ98945 standard; RNA; 21 BP
   Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                              20-OCT-2003; 2003US-00690276
                                                                                                                                                                                         28-JUL-2005 (first entry)
                                                                                                                                      Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                US2005112118-A1
                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                      02-DEC-1999;
                                                                                                                                                                                                                                                       26-MAY-2005
                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                  ADZ98945;
                                                                                                                                   Query Match
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The invention relates to a method for modulating, in a host cell, a protein-protein inceraction between a first protein which is PRAK (P38-corotein-protein inceraction between a first protein which is PRAK (P38-corotein-protein kinase or MAPKAPK5) and a second protein comparises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host protein-protein interaction between a first protein which is PRAK and a second protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, myositis, Crohn's disease, gastritis, colitis, colitis, ulcerative colitis, inflammatory bease, systemic lupus erythematosus, rhinitis, colitis, seleritis, chronic inflammatory polymouropathy, Tertiary Lyme disease, psoriasis, dermatis or eczema. The present sequence represents an siRNA (short interfering RNA) oligonucleotide targeting the continction in the seemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPKS) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein interaction; short interfering RNA; siRNA; RNA interference;
gene silencing; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human KU70 transcript siRNA sense oligonucleotide siRNA2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 7 A; 3 C; 4 G; 2 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 49; 296pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-00035344.
2002US-00099924.
2002US-00100503.
                                         2000US-00727384.
2000US-0255063P.
                                                                                                                                     2001US-0259572P.
                                                                                                                                                                                                                                2001US-00014814.
2001US-00024599.
2002US-00035343.
                                                                                          2000US-0256986P
                                                                                                                2001US-0259571P
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                                                                                                                                                                                                                                                                                                                                                                                                   (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADZ98947 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                               Heichman K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2005-371623/38.
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Synthetic.
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                                                                                                                                                           15-MAR-2001;
                                                                                       21-DEC-2000;
04-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                    18-MAR-2002;
                                                                    14-DEC-2000;
                                                                                                                                                                                                            23-JUL-2001;
                                                                                                                                                                                                                                                                                   04-JAN-2002;
                                                                                                                                                                                                                                                                                                      04-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Cimbora D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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ID ADZ9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
                                                        This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid bools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture of 10-50, particularly 15-25, T residues. The capture of 10-50, particularly 15-25, T residues. The capture comprises a sequence of (b) oncloid surface and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface to stranded nucleic acids to the surface is (quasi) covalent, supramolecular, is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucceal, cacids in a cells. It and matrix of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid pool, since the method is based on comparison of nucleic acid pools not individual cacuture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human KU70 transcript siRNA sense oligonucleotide siRNAl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/note= "2'-deoxythymine overhang"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. No. 8.1 vative 0; Mismatches
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Gaps

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BP.

Tue Nov

protein interaction; short interfering RNA; siRNA; RNA interference; gene silencing; ds.

/\*tag= a /note= "2'-deoxythymine overhang"

US2005112118-A1.

Location/Qualifiers

misc\_feature

Homo sapiens.

Synthetic

Human KU70 transcript siRNA sense oligonucleotide siRNA3

(first entry)

28-JUL-2005

ADZ98949 standard; RNA; 21 BP

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Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPK5) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
                                                                                                                                                                                                                                                                           Cimbora D, Heichman K, Bartel P, Mauck K, Bush A;
               20. .21
/*tag= a
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        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                            Disclosure, Fig 49; 296pp; English.
                                                                                                            99US-0168379P.
2000US-0185056P.
                                                                                                                                     2000US-0255063P.
2000US-0256986P.
                                                                                   20-OCT-2003; 2003US-00690276
                                                                                                    99US-0168377P
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2001US-0307233P.
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2001US-00024599
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                                                  US2005112118-A1.
                                                                                                                                                              04-JAN-2001;
15-MAR-2001;
19-MAR-2001;
23-JUL-2001;
                misc_feature
                                                                                                                                                                                                14-DEC-2001;
21-DEC-2001;
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                                                                                                                                             21-DEC-2000;
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                                                                  26-MAY-2005
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The invention relates to a method for modulating, in a host cell, a protein-protein interaction between a first protein which is PRAK (P38-cegulated/activated protein kinase or MAPRAPS) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein-protein interaction. The method is useful in modulating in a host cell a protein-protein interaction between a first protein which is PRAK and a second protein which is ERK3 for treating inflammation or inflammatory disorders, e.g., asthma, rheumatoid arthritis, modulating, invenile chronic arthritis, mostils Crohn's disease, proctitis, colitis, ulcarative colitis, inflammatory bowel disease, proctitis, pelvic inflammatory disease, systemic lupus erythematosus, rhinitis, collitis, coll
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0.7%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 8.1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
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The invention relates to a method for modulating, in a host cell, a protein-protein interaction between a first protein which is PRAK (P38-cegulated/activated protein kinase or MAPKARKS) and a second protein which is ERK3 (extracellular signal-regulated kinase 3). The method which is ERK3 (extracellular signal-regulated kinase 3). The method comprises administering to the cell a compound capable of modulating the protein interaction. The method is useful in modulating in a host cell a protein-protein interaction between a first protein which is PRAK inflammatory disorders, e.g., asthma, rheumatoid arthritis, juvenile chronic arthritis, myositis, cabhma, rheumatoid arthritis, juvenile chronic arthritis, myositis, crohn's disease, gastritis, colitis, ulcerative colitis, inflammatory bowel disease, proctitis, pelvic conjunctivitis, scleritis, chronic inflammatory polyneuropathy. Tertiary Lyme disease, psoriasis dermatitis or eczema. The present sequence copresents an siRNA (short interfering RNA) oligonucleotide targeting the KV70 transcript, which is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modulating, in a host cell, a protein-protein interaction between first protein, PRAK, (MAPKAPKS) and second protein, ERK3, (extracellular signal-regulated kinase 3) by administering modulating compound.
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Gaps ö

2038 GATGAAGGCTATCGTTGAG 2056

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||:||||||:||||:||:|||:||||| GAUGAAGGCUAUCGUUGAG 19

RESULT 888 ADZ98949

Bush A;

Χ, Mauck

Cimbora D, Heichman K, Bartel P,

WPI; 2005-371623/38.

2002US-00035344 2002US-00035343

04-JAN-2002;

18-MAR-2002; 2002US-00100503 (MYRI-) MYRIAD GENETICS INC.

01-DEC-2000; 2000US-00727384. 14-DEC-2000; 2000US-0255063P. 12-DEC-2000; 2000US-0255966P. 04-JAN-2001; 2001US-025971P. 04-JAN-2001; 2001US-0259772P. 15-MAR-2001; 2001US-02776179P. 19-MAR-2001; 2001US-0377613P. 14-DEC-2001; 2001US-0307233P. 14-DEC-2001; 2001US-0307233P. 21-DEC-2001; 2001US-0307239P.

99US-0168377P. 99US-0168379P. 2000US-0185056P.

02-DEC-1999; 25-FEB-2000;

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                                                   Gaps
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/note= "3' overhang comprising two 2'-deoxythymine
residues linked by a 5'-3' phosphodiester linkage"
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                                                                                                                                                                                                                                                                         Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive;
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                        0.7%; Score 19; DB 1; Length 21; 73.7%; Pred. No. 8.1e+02; tive 5; Mismatches 0; Indels
C; 3 G; 2 T; 5 U; 0 Other;
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                                                                             2059 GCTTCGCTTCACATACAGA 2077
                                                                                          2001US-0343818P.
2002US-00035344.
2002US-0346384P.
                                                                                                                                                                     AED42746 standard; RNA; 21 BP.
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2001US-0277013P.
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                                                   Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified; Synthetic.
Sequence 21 BP; 5 A; 6
                                     Best Local Similarity
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25-OCT-2001;
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07-JAN-2002;
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The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method of selecting modulators of the protein complex; a method of selecting modulators of the protein complex; a method of selecting modulators of an interaction between a first protein and a second protein complex as an analyor preventing of diseases and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders include cancer, adenomatous polyposis, isofamena, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, seteleptorosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an siRNA oligo which targets the transcript of a protein forming part of a protein-protein complex of the invention.
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                                                                                                                                  New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
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4.tag= a
//ncte= "3' overhang comprising two 2'-deoxythymine
residues linked by a 5'-3' phosphodiester linkage"
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                                                      Heichman K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 19; DB 1; Length 21; Pred. No. 8.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                      Wettstein DA,
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                                                          Sugiyama J,
                                                                                                                                                                                                                         Disclosure; Fig 62; 198pp; English.
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Local Similarity 73.7%;
les 14; Conservative
                 (MYRI-) MYRIAD GENETICS INC
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                                                        Cimbora D,
                                                                                             WPI; 2005-664172/68.
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                                                          Bartel P,
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Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive; Antidiabetic; Cardiant; Neuroprotective; Antiasthmatic; Antiinflammatory; Antibacterial; Osteopethic; Anorectic; Virucide, Anti-HIV; Nephrotropic; Antiarteriosclerotic; Muscular-Gen.; protein interaction; protein microarray; cancer; familial adenomatous polyposis; protein microarray; cancer; familial adenomatous polyposis; gastrointestinal-gen.; ischemia; cerebrovascular ischemia; autoimmune disease; dastemia; cerebrovascular ischemia; asthma; inflammation; sepsis; osteoporosis; obesity; viral infection; acquired immune deficiency syndrome; glomerulonephritis; atherosclerosis; muscular dystrophy; ss; short interfering RNA; RNA interference;
                                                                                                                                                                                       Protein interacting gene transcript siRNA antisense oligo #173
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15 - MAR - 2001; 2001US - 0259573F.
15 - MAR - 2001; 2001US - 0276259F.
15 - MAR - 2001; 2001US - 0276179F.
15 - APR - 2001; 2001US - 027403F.
17 - APR - 2001; 2001US - 028420F.
17 - APR - 2001; 2001US - 028420F.
17 - APR - 2001; 2001US - 028420F.
19 - APR - 2001; 2001US - 0287513F.
10 - UUL - 2001; 2001US - 0287513F.
22 - UUL - 2001; 2001US - 0347829F.
25 - OCT - 2001; 2001US - 0347829F.
25 - OCT - 2001; 2001US - 0343818F.
04 - JAN - 2002; 2002US - 0349843F.
17 - JAN - 2002; 2002US - 0349843F.
17 - MAR - 2002; 2002US - 0354899F.
18 - MAR - 2002; 2002US - 00099924.
      AED42749 standard; RNA; 21 BP.
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17-APR-2002; 2002US-00124550
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                                                                  AED42749;
      X X X Y Y Y X Y X X X X X X X X X X X Y X 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
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2001US-0276259P.
2001US-0276179P.
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2001US-0287513P.
2001US-0304101P.
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2001US-0284095P.
2001US-0284220P.
                                                        07-MAR-2005; 2005US-00075234
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Best Local Similarity 78.37
Best Local Similarity 78.37
Best Local Similarity
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30-APR-2001;
10-JUL-2001;
23-JUL-2001;
22-OCT-2001;
25-OCT-2001;
                                                                                                                                                                                                                                         19-MAR-2001;
16-APR-2001;
17-APR-2001;
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17-JAN-2002;
06-FEB-2002;
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14-MAR-2002;
18-MAR-2002;
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06-OCT-2005
                                                                                                                                                                                                             15-MAR-2001
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20. .21
/tag= a
/note= "3' overhang comprising two 2'-deoxythymine
residues linked by a 5'-3' phosphodiester linkage"

Location/Qualifiers

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New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
                                                                                                   Bartel P, Cimbora D, Sugiyama J, Wettstein DA, Heichman K;
(MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                 WPI; 2005-664172/68.
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Gaps

.; 0

0; Indels

4; Mismatches

1632 AGATTATACTGGAGAAGA 1650

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AED42749/c RESULT 891

2002US-00124767 2002US-00135802 Disclosure; Fig 62; 198pp; English

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The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method for selecting modulators of the protein complex; a method of selecting conclusions of an interaction between a first protein and a second protein complexes and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders associated with the protein complexes. The diseases and disorders include cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, osteoporosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an siRNA oligo which targets the transcript of a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive; Antidiabetic; Cardiant; Neuroprotective; Antiasthmatic; Antiniflammatory; Antibacterial; Osteopathic; Anorectic; Virucide; Anti-HIV; Nephrotropic; Antiarteriosclerotic; Muscular-Gen.; protein interaction; protein microarray; cancer; familial adenomatous polyposis; gastrointestinal-gen.; ischemia; cerebrovascular ischemia; autoimmune disease; darate disease; neurodegenerative disease; asthma; inflammation; sepsis; osteoporosis; obesity; viral infection; acquired immune deficiency syndrome; glomerulonephritis; atherosclerosis; muscular dystrophy; ss; short interfering RNA; RNA interference;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                               protein forming part of a protein-protein complex of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "3' overhang comprising two 2'-deoxythymine residues linked by a 5'-3' phosphodiester linkage"
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                                                                                                                                                                                                                                                                                                                                                                                        Query Match

0.7%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 8.1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                       Seguence 21 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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2001US-0259573P.
2001US-0276259P.
2001US-0277013P.
2001US-0284095P.
2001US-0284220P.
2001US-02844404P.
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/note= "3'
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The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises: a protein microarray comprising the protein complex; a method of selecting comparison of the protein complex; a method of selecting comparison of an interaction between a first protein and a second protein complex and the treating and/or preventing of diseases and disorders associated with the protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders include complexes, and in treating and/or preventing diseases and disorders include cancer, adenomatous polyposis, isolated, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, osteoporosis, obesity, viral infection, AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. This sequence represents an aiRNA oligo which targets the transcript of a coronary and protein-protein complex of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; Vasotropic; Cerebroprotective; Immunosuppressive; Antidiabetic; Cardiant; Neuroprotective; Antidathmatic; Antinflammatory; Antibacterial; Osteopathic; Anorectic; Virucide; Anti-HIV; Nephrotropic; Antiarteriosclerotic; Muscular-Gen.; protein interaction; protein microarray; cancer; familial adenomatous polyposis;
                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
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100.0%; Pred. No. 8.1e+02;
iive 0; Mismatches 0; Indels
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2002US-00035344.
2002US-0346384P.
2002US-0349843P.
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2002US-00098979.
2002US-00099924.
2002US-00100503.
               2001US-0304101P.
2001US-0307233P.
2001US-0347829P.
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2002US-00124767
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Best Local Similarity 100.
Matches 19, Conservative
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                                                                                                                            17-JAN-2002;
06-FEB-2002;
14-MAR-2002;
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07-JAN-2002;
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18-MAR-2002;
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                                                      22-OCT-2001;
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(MYRI-) MYRIAD GENETICS INC.
                                      WPI; 2005-664172/68.
                                     Bartel
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gastrointestinal-gen.; ischemia; cerebrovascular ischemia; autoimmune disease; diabetes; heart disease; neurodegenerative disease; asthma; inflammation; sepsis; osteoporosis; obesity; viral infection; acquired immune deficiency syndrome; glomerulonephritis; atherosclerosis; muscular dystrophy; ss; short interfering RNA; RNA interference;
                                                                                                                                                                                                                              /note= "3' overhang comprising two 2'-deoxythymine
residues linked by a 5'-3' phosphodiester linkage"
                                                                                                                                                                      Location/Qualifiers
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2001US - 0304233P
2001US - 0343818P
2002US - 00035344
2002US - 0346384P
2002US - 0346384P
2002US - 0354899P
2002US - 0354899P
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2001US-0284220P.
2001US-0284404P.
2001US-0285324P.
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/note= "3'
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                                                                                                                                 Unidentified; Synthetic.
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                                                                                               gene silencing
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15-MAR-2001;
19-MAR-2001;
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14-MAR-2002;
18-MAR-2002;
15-APR-2002;
17-APR-2002;
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25-OCT-2001;
04-JAN-2002;
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17-APR-2001;
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06-FEB-2002;
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The invention relates to a novel isolated protein complex having a first protein interacting with a second protein. The invention further comprises a protein microarray comprises, orderied microarray comprising the protein complex, a method for selecting modulators of the protein complex; a method of selecting modulators of an interaction between a first protein and a second protein; and the treating and/or preventing of diseases and disorders associated with the protein complexes. The protein complexes are useful in screening assays for identifying compounds effective in modulating the protein complexes, and in treating and/or preventing diseases and disorders associated with the protein complexes. The diseases and disorders cancer, adenomatous polyposis, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, neurodegenerative diseases, asthma, inflammatory disorders, sepsis, osteoporosis, obesity, viral infection,
New isolated protein complex having a first protein interacting with a second protein, useful for treating or preventing, e.g. cancer, ischemia, stroke, autoimmune diseases, diabetes, coronary heart disease, or asthma.
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Heichman K;

Wettstein DA,

Sugiyama J,

Cimbora D,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vaccine, cytostatic, virucidal; bactericidal, fungicidal, anti-parasitic, immunostimulatory, tumour; viral infection, bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                  of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                 Gaps
AIDS, glomerulonephritis, atherosclerosis, and muscular dystrophy. sequence represents an siRNA oligo which targets the transcript of protein forming part of a protein-protein complex of the invention.
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                                                                                                        DB 1; Length 21;
8.1e+02;
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                                                                      Seguence 21 BP; 5 A; 4 C; 2 G; 2 T; 8 U; 0 Other;
                                                                                                                 100.0%; Preq. ...
                                                                                                          Score 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory nucleic acid #52.
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                                                                                                                                                                                                   19 AAAGTGTACATCAGTAA 1
                                                                                                                                                                                                                                                                                                         AAF98936 standard; DNA; 22 BP.
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                                                                                                                                             19; Conservative
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                                                                                                                            Similarity
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                                                                                                          Query Match
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The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                  Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating non-allergic inflammatory diseases, such as psoriasis, eczialergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
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Pred. No. 8.5e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunostimulatory nucleic acid #52.
                                                                                                                                                               Immunostimulatory nucleic acid #55
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                                       ACD99369 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB36438 standard; DNA; 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KRIE/) KRIEG A M. (BERG/) BERG D J.
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                                                                                                                        25-SEP-2003
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                                                                                ACD99369;
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Best Local S
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Matches
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                     ACD99369,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lession, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osber-Webber Syndrome, myocardial angiogenesis, plaque wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inhibiting anglogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                          tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy, retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; oller.Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
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                                                                                                                                                                                                                                                                                                                                                             Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 18.8; DB 1; Length 22; 90.9%; Pred. No. 8.5e+02;
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    Indels
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0; Mismatches 2
                                                                                                                                                                                                                                                                                                                 Angiogenesis inhibitory oligonucleotide #61.
0; Mismatches
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                                           scleroderma; hypertrophic scar.
                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (COLE-) COLEY PHARM GROUP INC.
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                                                                                                                                                                                       ABS77577 standard; DNA; 22
                                                                                                                                                                                                                                                                            (first entry)
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    Conservative
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Best Local Similarity
Matches 20; Conserv
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RESULT 899
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                                                                                                                                                                                                      The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating,
                                                                                                                                                       Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic entil; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
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                                                                                                                                                                                                                                                                                              0.7%; Score 18.8; DB 1; Length 22; 90.9%; Pred. No. 8.5e+02; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                          Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
                                                                                                                   Petersen DM, Fouron Y;
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                                                                                                                                                                                     Claim 10; Page 6; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Non-CpG DNA oligonucleotide 37.
                                                                                                                                                                                                                                                                                                                                                                                                       ADG76036 standard; DNA; 22 BP
                                       02-FEB-2001; 2001US-00776479.
                                                         03-FEB-2000; 2000US-0179991P.
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                                                                                                                                                                                                                                                                                                                20; Conservative
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                                                                           (BRAT/) BRATZLER R L.
(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          renal cell carcinoma
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  US2003087848-A1
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                                                                                                                 Bratzler RL,
                   08-MAY-2003
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This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacyciol dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including carrinoma. This oligonucleotide sequence is a non-CpG DNA oligo of the
preventing or ameliorating immune system disorder or tumoral disease e.g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-cell; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Raposi's sarcoma;
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Pred. No. 8.5e+02;
0; Mismatches 2; Indels
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                                                                                                         Example 17; Page 81; 139pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22 AAAAACAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  melanoma.
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oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune seaponse is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipida. As such, these DNA oligos can be used in gene menliorating an immune system discorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is a non-CpG DNA oligo of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  se; antiasthmatic; antiallergic; dermatological; antiinflammatory;
antibacterial; virucide; immunoglobulin E antagonist; allergy;
immunostimulator; asthma; rhinitis; urticaria; dermatitis;
bacterial infection; viral infection.
                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                0.7%; Score 18.8; DB 1; Length 22; 90.9%; Pred. No. 8.5e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                              Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Allergic response suppressor oligonucleotide #61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 61; 235pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fouron Y;
                                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAAAA 2730
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02-FEB-2001; 2001US-00776479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-APR-2004; 2004US-00831778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADU89377 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 90.9
Matches 20; Conservative
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Э. Е.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BRAT/) BRATZLER
(PETE/) PETERSEN
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2004235774-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-FEB-2005
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                                                                                                                                                                                                                                              invention.
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(FOUR/)
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                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; immune stimulation; inflammation; psoriasis; inflammatory bowel disease; Crohns disease; ulcerative colitis; eczema; skin allergy; contact dermatitis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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                                                                                                                                         Gaps
     and viral
medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the method of the invention.
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llarity 90.9%; Pred. No. 8.5e+02;
Conservative 0; Mismatches 2; Indels
                                                                                                                                       2; Indels
                                                                                                       Length
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                                                                  Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
                                                                                                       DB 1;
                                                                                                     0.7%; Score 18.8; DB 1;
90.9%; Pred. No. 8.5e+02;
iive 0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory oligonucleotide, SEQ ID 55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 55; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               contact dermatitis or latex dermatitis.
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29-MAR-2002; 2002US-00112653.
                                                                                                                                                                                                                                                                                            AED74922 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                               12-JAN-2006 (first entry)
                                                                                                                         Local Similarity 90.9
nes 20; Conservative
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es 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                            Ouery Match
                                                                                                                                            Matches
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The invention relates to the use of one or more splice site-specific primers and optionally one or more arbitrary fragment length polymorphism (AFLP) primers, in a method for analyzing or amplifying a nucleic acid sequence or in the development of ECR primers. The invention also relates to using a PCR primer obtained by the method in the development of an assay, preferably for the analysis of splice sites. The method is useful in analyzing nucleic acid sequences for the presence or absence of splice site-associated polymorphisms. The splice site-specific primers and AFLP primers are useful in genotyping, genetic mapping, genetic profiling and DNA-identification techniques, e.g., to identify a specific species, subspecies, variety, race or individual, to establish the presence or basence of a specific inheritable trait and/or of a gene, or to determine the state of a disease. The splice site-specific primers allow for efficient/exponential amplification. The splice site-specific primers allow for efficient/exponential amplification. The splice site-specific findersprints. The primers have the capacity to target a large population of splice sites present in the genome. This sequence represents a splice-content of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of splice site-specific primers or combination of splice site-
specific primers and arbitrary fragment length polymorphism primer, for
analyzing/amplifying nucleic acid sequence, or in development of PCR-
                                                                                                                                                                        Arbitrary fragment length polymorphism; AFLP; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hogers RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Van Eijk MJT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; SEQ ID NO 12; 53pp; English.
                                                                                                                                    Splice site-specific primer #12
                                                                                                                                                                                               gene targeting; DNA mapping; ss
                      BP.
                                                                                                                                                                                                                                                                                                                                                     02-JUL-2004; 2004WO-NL000471.
                                                                                                                                                                                                                                                                                                                                                                                              02-JUL-2003; 2003WO-NL000486.
                  ADV92331 standard; DNA; 23
                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vogelaar A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-081957/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                  (KEYG-) KEYGENE NV
                                                                                                                                                                                                                                                                        WO2005003393-A2
                                                                                               07-APR-2005
                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                13-JAN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dirks RHG,
                                                        ADV92331;
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ADV92331,
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ö Gaps ö DB 1; Length 23; 2; Indels Score 18.8; DB 1; Pred. No. 8.7e+02; 0; Mismatches 2 0.7%; Ouery Match
Best Local Similarity 90...
Best Local Similarity 50...

ઠે 유 RESULT 903 ADC17041/c ID ADC17041 standard; DNA; 51 BP. ADC17041; a x 2

(first entry)

Human single nucleotide polymorphism (SNP) region Seq ID143.

sequence polymorphism analysis; human identity; human relatedness; single nucleotide polymorphism; SNP; genetic disease; cytostatic; immunosuppressive; antiinflammatory; neuroprotective; antimicrobial; fatty acid metabolism; glycolysis; amino acid metabolism; paternity analysis; forensic; autoimmune disease; cancer; nervous system; infection; pathogenic microorganism; human; ds.

Homo sapiens

Location/Qualifiers replace (26,G) variation

\*tag=

/standard\_name= "Single nucleotide polymorphism"

WO200029622-A2.

25-MAY-2000.

99WO-US027283, 17-NOV-1999;

98US-0109024P. 17-NOV-1998; 16-NOV-1999;

(CURA-) CURAGEN CORP.

Shimkets RA,

WPI; 2000-399731/34. P-PSDB; ADC16824.

for

Novel polynucleotide and polypeptide including one or more single nucleotide polymorphisms, useful for diagnosing and treating conditions associated with the presence of sequence polymorphism in humans and animals.

Claim 1; SEQ ID NO 143; 187pp; English.

comprise 217 defined polymorphic sequences. Sequence polymorphism-based analysis of nucleic acid sequences can augment or replace previously known methods for determining the identity and relatedness of individuals. Single nucleotide polymorphisms (SNPs) tend to occur with great frequency throughout the genome and may be located close to loci of interest. Such variations cause or be closely linked to pathological conditions (genetic diseases). Hence the SNPs of the invention may be useful in the development of compounds with cytostatic, imunosuppressive, antihinflammatory, neuroprotective or antimicrobial activities. Regulators of metabolic pathways such as fatty acid metabolism, glycolysis, and amino acid metabolism may also be developed. The compounds may be useful for treating a subject suffering from or at risk for a pathology associated with the presence of a sequence polymorphism. SNP detection is also useful in paternity analysis and correction is also useful in paternity analysis and correction is also useful in paternity analysis and contract application. Polymorphisms may contribute to the phenotype of an order of a pathology associated with the presence of a sequence of an order of the phenotype of the phen organism and phenotypic traits include genetic diseases such as autoimmune diseases, cancer, diseases of the nervous system and infection by pathogenic microorganisms. The present sequence is the sequence surrounding and including a human SNP of the invention. This invention relates to novel isolated nucleotide sequences which 

Sequence 51 BP; 22 A; 6 C; 14 G; 9 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 18.8; DB 1; Length 51; 33.0%; Pred. No. 1e+03; ; \_\_; 1e+03; .hes 17; Indels 0; Mismatches ilarity 63.0%; Conservative Local Similarity 29; Query Match Best Loc Matches

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The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcribtome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The coligonucleotide libraries are useful for detecting mRNAs from a coligonucleotide libraries are useful for detecting mRNAs from a coligonucleotide libraries are useful for detecting mRNAs from a coligonucleotide libraries may also be used as specialised minic detecting RNA transcripts of a sub-transcriptome under a particular colibraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue colipraries to detect transcripts of a sub-transcriptome under a particular coliponental specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect coliprariants of a transcriptome of a patient suffering from a particular disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from trans. In the printed specification, but was obtained in electronic format condition from the printed specification, but was obtained in electronic format colifectly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit of genome, useful for detecting tissue-, pathology-, and developmental-
                                                                                                                             Human spliced transcript detection oligonucleotide SEQ ID NO:31969.
                                                                                                                                                                     mouse; rat; splice transcript; detection; RNA transcript; variant; transcriptome; oligonucleotide library; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 0.7%; Score 18.6; DB 1; Length 60; Local Similarity 57.9%; Pred. No. 9.7e+02; nes 33; Conservative 0; Mismatches 24; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; SEQ ID NO 31969; 47pp; English.
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ABN59221 standard; DNA; 60 BP.
                                                                                                                                                                                                                                                                                                                                                                   20-JUL-2001; 2001WO-IB001903
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02-MAY-2001; 2001US-0287724P.
                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-257383/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific genes.
                                                                                                                                                                                                                                                                              WO200210449-A2
                                                                                                                                                                         Human; mouse;
                                                                                                                                                                                                                                        Homo sapiens.
                                                                                    15-JUL-2002
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                                          ABN59221;
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of of
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                                                                                                                                                                                                                                                                                                                                                                                                                                          A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GRNESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gapa
                                                                                                              Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                   Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                      by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 5; 11pp; Japanese.
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AAQ75569 standard; DNA; 20
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Best Local Similarity 95.0
Matches 19; Conservative
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                              AAQ75569;
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alysis of cDNA and gene expression - by amplification of mRNA followed digestion with restriction enzymes.
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                         0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ive 0; Mismatches 1; Indels
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AAQ75579 standard; DNA; 20
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by digestion with restriction enzymes.
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A method for the analysis of cDNA comprises (a) preparing an aggregate c double-stranded cDNAs by using an aggregate of mRNAs and a plural type c labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                          by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                           Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                          2706 ACTAAAAAAAAAAAAAA 2725
                                                                                             aggregate; restriction enzyme; ss.
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          BP.
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       AAQ75593 standard; DNA; 20
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                                             (first entry)
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                                                                                                                                                                                                                             WPI; 1995-018287/03.
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Matches 19; Conserv
                                                                                                                                JP06303997-A.
                                             04-AUG-1995
                                                                                                                                                                    16-APR-1993;
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                                                                                                             Synthetic.
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                           AAQ75593,
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AAQ75593/c
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of
                                                                                                                                        Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                           A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AA(75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 18.4; DB 1; Length 20;
95.0%; Pred. No. 8.6e+02;
ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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ID AAQ75601 standard; DNA; 20
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                                                                  WPI; 1995-018287/03.
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Matches 19; Conserv
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                             JP06303997-A.
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          Synthetic.
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Matches
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AAQ75595/
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of
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                    Gaps
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                                                                                                                                                                       Reverse transcription primer used in cDNA analysis technique.
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Pred. No. 8.6e+02;
0; Mismatches 1; Indels
 Length 20;
                    Indels
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 0.7%; Score 18.4; DB 1;
95.0%; Pred. No. 8.6e+02;
iive 0; Mismatches 1;
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                                       2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                            AAQ75581 standard; DNA; 20 BP
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Conservative
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Query Match
Best Local Similarity 95.0
Matches 19; Conservative
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Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                     by amplification of mRNA followed
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Pred. No. 8.6e+02;
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                                                                                                                                                                                                                           expression -
                                                                                                                                                                                                                                                            by digestion with restriction enzymes.
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                                                                                                                                                                                                                              of cDNA and
                                                                                                                                                                      WPI; 1995-018287/03.
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double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New stem cell factor polypeptide(s) - for stimulating the growth of primitive progenitor cells, esp. for treating disorders involving blood
                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalian stem cell factor (SCF) cDNA oligonucleotide primer 220-3.
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thrombocytopenia; leucopenia; AIDS; immunodeficiency; bone graft;
transplant; neoplasia; myelosuppression; bone marrow; ss.
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                                                                                                                                                                                  Score 18.4; DB 1; Length 20;
Pred. No. 8.6e+02;
0; Mismatches 1; Indels
                                                                                                                                             Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
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90US-00537198.
90US-00573616.
90WO-US005548.
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(first entry)
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Best Local Similarity 95.0°
Matches 19; Conservative
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15-MAY-1996
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28-SEP-1990;
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amulatication and sequencing of mammalian stem cell factor (SCF). Non-
naturally occuring SCF and C-terminally truncated polypeptides, having
amino acid sequences sufficiently duplicative of naturally courring SCF,
stimulate growth of primitive progenitors such as haematopoietic
progenitor cells, neural stem cells and primordial germ stem cells. The
peptides can be used in a composition for treating leucopenia, anaemia or
thrombocytopenia, for enhancing engraftment of bone marrow during
transplantation or for bone marrow recovery after chemotherapy or
cradiation-induced bone marrow aplasia or myelosuppression. They can also
be used for treating neoplasia, nerve damage, infertility, intestinal
camage or myeloproliferative disorders. Antibodies may be raised against
the peptides for use in detection or neutralisation of SCF in serum. SCF
may be useful for the treatment of AIDS and severe combined
immunodeficiency (SCID) states alone or in combination with other factors
such as IL-7. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                ö
may be useful for the treatment of AIDS and severe combined immunodeficiency (SCID) states alone or in combination with other factors such as IL-7. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New stem cell factor polypeptide(s) - for stimulating the growth of primitive progenitor cells, esp. for treating disorders involving blood
                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalian stem cell factor (SCF) cDNA oligonucleotide primer 220-11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stem cell factor; progenitor; haematopoiesis; SCF; anaemia;
thrombocytopenia; leucopenia; AIDS; immunodeficiency; bone graft;
transplant; neoplasia; myelosuppression; bone marrow; ss.
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0
                                                                                                      0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ve 0; Mismatches 1; Indels
                                                                         Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                2707 CTAAAAAAAAAAAAAAA 2726
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90US-00537198.
90US-00573616.
90WO-US005548.
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                                                                                                                           Local Similarity 95.0
nes 19; Conservative
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                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                cell factor; SCF; haematopoletic progenitor cell; blood forming;
                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                               primitive progenitor cell; haematopoletic disorder; syngeneic; allogeneic; aucologous bone marrow transplant; gene therapy; transfection; haematopoletic stem cell; acute blood loss; neoplasia;
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0
                                 Score 18.4; DB 1; Length 20;
Pred, No. 8.6e+02;
                                                                       1; Indels
Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                Stem cell factor universal oligonucleotide 220-3.
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                                                                        0; Mismatches
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                                                                                                            2707 CTAAAAAAAAAAAAAAAA 2726
                                                                                                                                              20 CGAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                        BP.
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90US-00573616.
90WO-US005548.
                                     0.7%;
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                                                                                                                                                                                                                                        AAA13752 standard; DNA; 20
                                                                                                                                                                                                                                                                                                             (first entry)
                                   Query Match 0.7
Best Local Similarity 95.0
Matches 19, Conservative
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28-SEP-1990;
01-OCT-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                           AAA13752;
                                                                                                                                                                                                    RESULT 920
                                                                                                                                                                                                                                                                                                                                                                                      Stem
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A method has been developed of making haematopoietic cells suitable for administration to a subject. The method comprises: (a) obtaining cadministration to a subject. The method comprises: (a) obtaining be administration to a subject. The method comprises: (a) obtaining the cells confirmation and one or more of the biological properties of naturally confirmation and one or more of the biological properties of naturally confirmation and one or more of the biological properties of naturally confirmation and one or more of the biological properties of naturally confirmation and cells including early haematopoietic progenitor cells which are capable of maturing early haematopoietic progenitor cells which are capable of maturing to erythroid, megakaryocyte, confirmation to properties cells of both myeloid and lymphoid lineages of increases in haematopoietic cells of both myeloid and lymphoid lineages. SCF is useful for treating haematopoietic disorders. The method is useful for expanding early haematopoietic progenitors in syngenetc, allogeneic confirmation bene maturow transplant. SCF is useful for enhancing the cefliciency of gene therapy based on transfecting haematopoietic stem cells. SCF is also useful for combating the myelosuppressive effects of anti-HIV drugs such as AZT and for enhancing haematopoietic recovery
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Production of hematopoietic cells suitable for administration to a subject using progenitor cells and expanding the cells using stem cell
oligonucleotide which is used in an example from the present invention
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                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stem cell factor; SCF; haematopoietic progenitor cell; blood forming primitive progenitor cell; haematopoietic disorder; syngeneic; allogeneic; autologous bone marrow transplant; gene therapy; transfection; haematopoietic stem cell; acute blood loss; neoplasia;
                                                                                                                                                                                           ö
                                                                                                                              Length 20;
                                                                                                                                                                                        1; Indels
                                                          Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stem cell factor universal oligonucleotide 220-11.
                                                                                                                                 DB 1;
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                                                                                                                           Score 18.4; DB 1;
Pred. No. 8.6e+02;
0; Mismatches 1;
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90US-00537198.
90US-00573616.
90WO-US005548.
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                                                                                                                                 0.7%;
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90EP-00310899
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                                                                                                                                                                                                  Conservative
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                                                                                                                                                                 Local Similarity
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28-SEP-1990;
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04-OCT-1990;
                                                                                                                                                                                                  19;
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                                                                                                                                    Query Match
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Gaps

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1; Indels

Pred. No. 8.6e+02; 0; Mismatches 1

95.0%;

19; Conservative

Best Local Similarity Matches 19; Conserv

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Enhancing efficiency of transfer of polynucleotide into a target mammalian cell in vitro, involves exposing cell that expresses a stem cell factor receptor to stem cell factor, and introducing polynucleotide into cell in vitro.
after acute blood loss and as a boost to the immune system for fighting neoplasia (cancer). The present sequence represents a universal oligonuclectide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                         Stem cell factor; SCF; stem cell factor receptor; blood cell disorder; gene therapy; PCR primer; mutagenesis; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                           Universal stem cell factor (SCP) related oligonucleotide SEQ ID NO:32.
                                                                                                                                      Gaps
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                                                                                                  0.7%; Score 18.4; DB 1; Length 20;
95.0%; Pred. No. 8.6e+02;
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                                                               Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                                 Pred. No. 8.66
0; Mismatches
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90US-00537198.
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90US-00982255.
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95US-00446553.
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                                                                                                                Local Similarity 95.0
les 19; Conservative
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24-AUG-1990
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0.7%; Score 18.4; DB 1; Length 20;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enhancing efficiency of transfer of polynucleotide into a target mammalian cell in vitro, involves exposing cell that expresses a stem cell factor receptor to stem cell factor, and introducing polynucleotide
                                                                                                                                                                                                                         Stem cell factor; SCF; stem cell factor receptor; blood cell disorder; gene therapy; PCR primer; mutagenesis; probe; ss.
                                                                                                                                                                                             Universal stem cell factor (SCF) related oligonucleotide SEQ ID NO:34.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Martin FH;
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2707 CTAAAAAAAAAAAAAA 2726
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920S-00982255.
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980S-00005893.
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90US-00573616
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                                                                                                     AAH41333 standard; DNA; 20
                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     into cell in vitro.
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                                                                                                                                                                                                                                                                                                    US6207454-B1.
                                                                                                                                                             21-AUG-2001
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24-AUG-1990;
01-OCT-1990;
25-NOV-1992;
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                                                                                                                                                                                                                                                                      Synthetic.
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                            20
                                                                                                                                   AAH41333;
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                                                                         RESULT 92
AAH41333/
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Tue Nov

ngs.res

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The present sequence for universal PCR primer 220-11 is 1 of 8 universal oligonucleotides (AASO4110-AASO4117) used in the isolation of the human SCF (Stem cell factor) CDNA sequence. The present invention relates to novel stem cell factors (AAU02454) AAU02460, AAU02461) and the collection of the novel stem cell factors (AAU02453-AAU02454). AAU02461) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells including early haematopoietic progenitor cells. The invention also describes SCF peptides (AAU02462-AAU02481) and the oligonucleotides (AASO4081-AASO4117) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in the isolation of human and rat SCF sequences. The polynucleotide metastatic carcinoma, acute leukaemia, multiple such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple cyeloma, Hodgkin's disease, lymphoma, Gaucher's disease, anaemia, congestive splenomegaly, Kala azar, sarcoidosis, military tuberculosis, disseminated fungus disease, Pulminating septicemia, malaria, vitamin B12 and folic acid deficiency, pyridoxine deficiency, and hypopigmentation disorders such as piebaldism and vitiligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acids encoding stem cell factor useful for treating disorders involving blood cells, e.g. leukemia, splenomegaly, Hodgkin's disease, Kala azar, anemia and septicemia.
                                                                                                                                                                                                    blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly; anaemia; Kala azar; septicemia; malaria; hypopigmentation disorder;
                                                                                                                                                                                    Human; stem cell factor; SCF; early haematopoietic progenitor cell;
                                                                                                                                                Human SCF (stem cell factor) cDNA universal PCR primer 220-11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suggs SV, Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Fig 12C; 209pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            90US-00537198.
90US-00573616.
90US-00589701.
                                  AAS04113 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                       95US-00482918
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                                                                                                              (first entry)
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Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ZSEB/) ZSEBO K M.
(BOSS/) BOSSELMAN R A.
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-298941/31.
                                                                                                                                                                                                                                                 PCR primer; 88
                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                     07-JUN-1995;
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                                                                                                              29-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                              16-OCT-1989;
                                                                                                                                                                                                                                                                                                                                                                 27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zsebo KM,
                                                                        AAS04113;
RESULT 924
                    AAS04113
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The present sequence for universal PCR primer 220-3 is 1 of 8 universal coligonucleotides (AAS04110-AAS04117) used in the isolation of the human coligonucleotides (AAU02453-AAU02458, AAU02460, AAU02461) and the covel stem cell factors (AAU02453-AAU02458, AAU02460, AAU02461) and the color stem cell factors (AAU02453-AAU02484) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells coloring carly haematopoietic progenitor cells. The invention also describes SCF peptides (AAU02462-AAU02481) and the oligonucleotides (AAS04117) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in gene therapy. It is useful for treating disorders involving Blood cells can melalogize disease, metastatic carcinoma, acute leukaemia, multiple congestive splenomegaly, Kala azar, sarcoidosis, military tuberculosis, congestive splenomegaly, Rala azar, sarcoidosis, military tuberculosis, alsementated fungus disease, hulminating septicemia, malaria, vitemin Bl2 coloring active carcinomes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acids encoding stem cell factor useful for treating disorders involving blood cells, e.g. leukemia, splenomegaly, Hodgkin's disease, Kala azar, anemia and septicemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and folic acid deficiency, pyridoxine deficiency, and hypopigmentation disorders such as piebaldism and vitiligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                            Human, stem cell factor; SCF; early haematopoietic progenitor cell;
blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly;
anaemia; Kala azar; septicemia; malaria; hypopigmentation disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                            Human SCF (stem cell factor) cDNA universal PCR primer 220-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 18.4; DB 1;
Pred. No. 8.6e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Suggs SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; Fig 12C; 209pp; English.
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90US-00537198.
90US-00573616.
90US-00589701.
93US-00172329.
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              AAS04111 standard; DNA; 20
                                                                               (first entry)
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nes 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zsebo KM, Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZSEBO K M.
BOSSELMAN R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-298941/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MART/) MARTIN F H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUGGS S
                                                                                                                                                                                                    PCR primer; ss.
                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                      US6207417-B1.
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24-AUG-1990;
01-OCT-1990;
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                                                 AAS04111;
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AAS04111
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Gaps

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2707 CTAAAAAAAAAAAAAA 2726 

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20

RESULT 926

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11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
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                                                                                                          09-NOV-1994;
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                                                                                     27-MAR-2001
                                                                                                                                                                       25-NOV-1992;
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                                                                                                                                                                                                                   Zsebo KM,
                                             Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH23889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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   g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides the protein and coding sequences of mammalian stem cell factors (SCFs). These are capable of stimulating the growth of early haematopoietic progenitor cells, neural stem cells and primordial germ stem cells. The sequences are useful in the treatment of leukaemias, haematopoietic disorders, aplastic anaemia, paroxysmal nocturnal haemoglobinuria, malaria, pigmentation disorders, neurological and intestinal damage, infertility, AIDS and severe combined immunodeficiency (SCID). The present sequence is primer used to amplify an SCF in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                               Novel isolated non-human mammalian stem cell factor polypeptide stimulating growth of early hematopoietic progenitor cells, useful for treating aplastic anemia, lymphoma, Letterer-Siwe disease, Kala azar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                          Human; rat; mammal; stem cell factor; SCF; cell growth stimulation; gene therapy; haematopoietic disorder; aplastic anaemia; leukaemia; neurological damage; intestinal damage; infertility; AIDS; SCID; severe combined immunodeficiency; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; rat; mammal; stem cell factor; SCF; cell growth stimulation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; tive 0; Mismatches 1; Indels
                                                                      Mammalian stem cell factor PCR primer SEQ ID NO: 32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalian stem cell factor PCR primer SEQ ID NO: 34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                        Martin FH;
                                                                                                                                                                                                                                                                                                                        Suggs SV,
                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Fig 12C; 209pp; English.
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90US-00573616.
90US-00589701.
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                                                                                                                                                                                                                                                                              92US-00982255
         AAF89091 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF89093 standard; DNA; 20
                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                       Zsebo KM, Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                           WPI; 2001-353108/37.
                                                                                                                                                                                                                                                                                                   (AMGE-) AMGEN INC.
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24-AUG-1990;
01-OCT-1990;
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                                                  13-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                 sarcoidosis.
                                                                                                                                                                                                                                                                              25-NOV-1992
                                                                                                                                                                                          27-MAR-2001
                                                                                                                                                                                                                                    16-OCT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUL-2001
                                                                                                                                                 Mammalia.
                             AAF89091;
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AAF89091/c
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The present invention provides the protein and coding sequences of mammalian stem cell factors (SCFs). These are capable of stimulating the growth of early haematopoietic progenitor cells, neural stem cells and primordial germ stem cells. The sequences are useful in the treatment of leukaemias, haematopoietic disorders, aplastic anaemia, paroxysmal nocturnal haemoglobinuria, malaria, pigmentation disorders, neurological and intestinal damage, infertility, AIDS and severe combined immunodeficiency (SCID). The present sequence is primer used to amplify an SCF in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated non-human mammalian stem cell factor polypeptide stimulating growth of early hematopoietic progenitor cells, useful for treating aplastic anemia, lymphoma, Letterer-Siwe disease, Kala azar, sarcoidosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
gene therapy; haematopoietic disorder; aplastic anaemia; leukaemia; neurological damage; intestinal damage; infertility; AIDS; SCID; severe combined immunodeficiency; PCR primer; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Fig 12C; 209pp; English.
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90US-00537198.
90US-00573616.
90US-00589701.
92US-00982255.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19; Conservative
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primer; ss.
                                                                                                                  (AMGE-) AMGEN
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11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
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                                                               10-APR-1991;
11-JUN-1990;
                     24-AUG-1990;
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                                                                                                                                                                                                                                                                                                                                  disease.
                                           01-OCT
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AAS04214/c
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                                                                                                                                                                                                                                                                                                                             New stem cell factor polypeptides and their analogs which stimulate growth of early hematopoietic progenitors, useful for treating aplastic anemia, carcinoma, multiple myeloma, vitiligo, kala azar, Hodgkin's
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blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly;
anaemia; Kala azar; septicaemia; malaria; hypopigmentation disorder;
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90US-00573616.
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                                                                                                                                                                                                                                                                                      WPI; 2001-256683/26.
                                                                                                                                                                                         (AMGE-) AMGEN INC.
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    25-NOV-1992;
                                                                                                                                        10-APR-1991;
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01-OCT-1990;
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The present sequence for universal PCR primer 220-11 is 1 of 8 universal coligonucleotides (AAH23888-AAH23895) used in the isolation of the human coligonucleotides (AAH23888-AAH23895) used in the isolation of the human correct form call factor) CDNA sequence. The present invention relates to novel stem cell factors (AAB73561-AAB73568, AAB73571-AAB73576) and the polynucleotides encoding them. SCF stimulate primitive progenitor cells including early heematopoietic progenitor cells. The invention also describes SCF peptides (AAB73578-AAB7359) and the oligonucleotides (AAH23899-AAH2389) used in the isolation of human and rat SCF sequences. The polynucleotide encoding SCF is useful for producing SCF and useful in SCF gene threapy. It is useful for treating disorders involving blood cells such as myelofibrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, anaemia, congestive applenomegaly, Kala azar, sarcoidosis, military tuberculosis, congestive planomegaly kala azar, sarcoidosis, military tuberculosis, consentinated fungus disease, Pulminating septicaemia, malaria, vitamin B12 and folic acid deficiency, pyridoxine deficiency, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                  New stem cell factor polypeptides and their analogs which stimulate growth of early hematopoietic progenitors, useful for treating aplastic anemia, carcinoma, multiple myeloma, vitiligo, kala azar, Hodgkin's
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90US-00537198.
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blood disorder; leukaemia; Hodgkin's disease; lymphoma; splenomegaly;
anaemia; Kala azar; septicemia; malaria; hypopigmentation disorder;
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                                                                                                                            Isolated DNA sequence, encoding polypeptide product useful for stimulating growth of early hematopoietic progenitor cells.
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Pred. No. 8.6e+02;
0; Mismatches 1; Indels
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                                                                  Martin FH;
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                                                                  Suggs SV,
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                                                             Bosselman RA,
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Best Local Similarity 95.0
Matches 19; Conservative
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                                (AMGE-) AMGEN INC.
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25-NOV-1992;
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                                                               Zsebo KM,
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P X Y X Y B B B B X B X B X S X X X X X X B X X 
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blood disorder, Hodgkin's disease, vitamin B12, folic acid deficiency;
hypopigmentation disorder, viral disorder, AIDS; PCR primer; ss.
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                            Isolated DNA sequence, encoding polypeptide product useful for
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                                          stimulating growth of early hematopoietic progenitor cells.
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                                                                     Example 3; Fig 12C; 167pp; English
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Best Local Similarity 95.05
Marches 19; Conservative
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SUGGS S V.
WPI; 2001-281051/29.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-APR-1991;
25-NOV-1992;
21-DEC-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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01-OCT-1990
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AAS10447/c
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                                                                                                     The present sequence for universal PCR primer 220-3 is 1 of 19 PCR primers (AAS10415-AAS10453) used to amplify various portions of the human SCF CNNA sequence. The sequence is described in an invention relating to novel stem cell factors, the polynucles encoding them and methods for producing the stem cell factors. The methods involve increasing the number of early haematopoietic progenitor cells in human peripheral blood polypeptide. The methods are useful for the treatment of blood disorders, including myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma (Gaucher's disease, Niemann-Pick disease, refractory anamia, malaria, vitamin Bl2 and folic acid deficiency, hypopigmentation disorders i.e. piebaldism and viral induced disorders, including AIDS
                        Increasing the number of early hematopoietic progenitor cells in the peripheral blood useful for the treatment of blood disorders including Hodgkin's disease comprises the administration of human stem cell factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; stem cell factor; SCF; haematopoietic progenitor cell;
blood disorder; Hodgkin's disease; vitamin B12; folic acid deficiency;
hypopigmentation disorder; viral disorder; AIDS; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human stem cell factor (SCF) cDNA universal PCR primer 220-11.
                                                                                                                                                                                                                                                                                                                              0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                                                                                      Seguence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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                                                                                 Example 3; Fig 12C; 210pp; English.
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90US-00537198.
90US-00573616.
90US-00684535.
92US-00684535.
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BOSSELMAN R A.
SUGGS S V.
MARTIN F H.
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WPI; 2001-407312/43.
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25-NOV-1992;
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Increasing the number of early hematopoietic progenitor cells in the

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peripheral blood useful for the treatment of blood disorders including Hodgkin's disease comprises the administration of human stem cell factor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel non-naturally-occurring stem cell factor polypeptide, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorders i.e. piebaldism and viral induced disorders, including AIDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18.4; DB 1;
Pred. No. 8.6e+02;
0; Mismatches 1;
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                                                                                                                           English.
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Best Local Similarity 95.0",
Best 19; Conservative
                                                                                                                               Example 3; Fig 12C; 210pp;
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(BOSS/) BOSSELMAN R A
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
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Example 3; Fig 12C; 217pp; English.
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                                                                                                       exemplification of the invention
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                                                                                                                            19; Conservative
                                                                                                                         Best Local Similarity
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                                                                                                                                                                   AAD35466;
                                                                                                                     Query Match
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                                                                                                                                                     RESULT 935
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The present invention relates to novel non-naturally-occurring stem cell factor (SCF) polypeptides having an amino acid sequence sufficiently courtains (SCF) polypeptides having an amino acid sequence sufficiently courtains (SCF) collogical activity of naturally occurring SCF. Sequences of the invention are useful for treating leucopaenia, thrombocytopaenia, canamia and for enhancing bone marrow recovery in treatment of radiation, engraftment of bone marrow during transplantation in mammals and chemical or chemotherapeutic induced bone marrow aplasia or myelosuppression. They are also useful for treating acquired immune deficiency in a human, nerved amage in a mammal. SCF sequences are useful for preparing biologically carise polymer polymer polymer polymer polymer polymer polymer polymer in the sequence are useful for preparing biologically active polymer polymer polymer cells with a gene, and transfer of a gene into a mammal. They are useful for treating myelofibrosis, myelosclerosis, costeopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, costeopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, corgestive splenomegaly, Kala azar, sarcoidosis, primary splenomegaly, splenomegaly, kala azar, sarcoidosis, primary splenomegaly, betterer-Siwe disease, refractory explenome, and folicancy vitamin Bloom disorders such as piebaldism, AlDS (acquired immune deficiency, vitamin Bloom of disorders such as piebaldism, AlDS (acquired immune deficiency syndrome) and liting or the present sequence is a PCR primer which is used for an english and an anticipal sequence is a poly primer.
                                                                                                                                                                                                                                                                                                                                                                                         Novel non-naturally-occurring stem cell factor polypeptide, useful for treating leucopenia, thrombocytopenia, anemia and for enhancing engraftment of bone marrow during transplantation in a mammal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.7%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 8.6e+02;
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                                                                                                                                                                                                                                                         Suggs SV, Martin FH;
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                 95US-00449653.
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                                                                                                                                                                                                                                                         Zsebo KM, Bosselman RA,
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                                                                          (ZSEB/) ZSEBO K M.
(BOSS/) BOSSELMAN R A.
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
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                 24-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                syndrome, congestive splenomegaly, Kala azar, sarcoidosis, primary splenic pancytopaenia, disseminated fungus disease, malaria, military tuberculosis, Pulminating septicaemia, pyridoxine deficiency, vitamin Bl2 and folic acid deficiency, Diamond Blackfan anaemia, hypopigmentation disorders such as piebaldism, AIDS (acquired immune deficiency syndrome) and vitiligo. The present sequence is a PCR primer which is used for amplifying the 5' end of rat SCF CDNA. This sequence is used in the
                                                                                                                                                                              present invention relates to novel non-naturally-occurring stem cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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       rrearing leucopenia, thrombocytopenia, anemia and for enhancing engraftment of bone marrow during transplantation in a mammal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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treating leucopenia, thrombocytopenia,
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Gaps

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1; Indels

Synthetic

98US-00005243.

12-JAN-1998; 14-FEB-2002

US2002018763-A1.

Rattus

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04-OCT-1990; 2002EP-00008587
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11-JUN-1990;
24-AUG-1990;
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04-OCT-1990
                      16-OCT-1989
                                    11-JUN-1990
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ADE52460/c
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                                                                                                                                                                                                                                                                                                                    polynucleotide sequences encoding the SCEs, and methods of producing them. SCFs are involved in the blood-forming (haematopoietic) system in mammals, particularly humans. The method of the invention is useful for the production of human SCF. The stem cell factors are useful to treat disorders involving blood cells e.g. metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, refractory erythroblastic anaemia, miliary tuberculosis, disseminated fungus disease, malaria, and vittiligo. The present sequence representing a universal oligonucleotide for SCF DNA is used in the examples of the
                                                                                                                                                                                                                           Production of a human stem cell factor (SCF) polypeptide for treating disorders involving blood cells, such as leukemia, comprises culturing mammalian cells comprising non-human SCF promoter DNA linked to DNA encoding the human SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stem cell factor; SCF; blood-forming system; blood cell disorder; haematopoietic system; metastatic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; malaria; vitiligo; refractory erythroblastic anaemia; miliary tuberculosis; cytostatic; aliseminated fungus disease; haematopoietic; tuberculostatic; antianaemic; antifungal; antimalarial; dermatological; ss.
                                                                                                                                                                                                                                                                                                            present invention relates to novel stem cell factors (SCFs),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

0.7%; Score 18.4; DB 1; Length 20;
Best Local Similarity 95.0%; Pred. No. 8.6e+02;
Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                    Zsebo KM, Suggs SV, Bosselman RA, Martin FH;
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                                                                   89US-00422383.
90US-00537198.
90US-00573616.
90WO-US005548.
90US-00589701.
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                                                                                                                            90EP-00310899
95EP-00105391
                                                2002EP-00008587
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                                                                                                                                                                                                          WPI; 2002-684093/74.
                                                                                                                                                               AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                              present invention
                                               04-OCT-1990;
   EP1241258-A2
                                                                                                                                       04-OCT-1990;
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                         18-SEP-2002
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The present invention relates to novel stem cell factors (SCFs), polynucleotide sequences encoding the SCFs, and methods of producing them. SCFs are involved in the blood-forming (haemaropoietic) system in mammals, particularly humans. The method of the invention is useful for the production of human SCF. The stem cell factors are useful to treat disorders involving blood cells e.g. metastatic carcinoma, acute erythroblastic anaemia, Hodghin's disease, lymphoma, refractory erythroblastic anaemia, miliary tuberculosis, disseminated fungus disease, malaria, and vitiligo. The present sequence representing a universal oligonucleotide for SCF DNA is used in the examples of the
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                                                                                                                                                                                                                                                                                                                                                                                                                           production of a human stem cell factor (SCF) polypeptide for treating disorders involving blood cells, such as leukemia, comprises culturing mammalian cells comprising non-human SCF promoter DNA linked to DNA encoding the human SCF.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              Martin FH;
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                                                                                                                                                                                                                                                                                                              Bosselman RA,
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90US-00537198.
90US-00573616.
                       90US-00537198.
90US-00573616.
90WO-US005548.
90US-00589701.
90EP-00310899.
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89US-00422383
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                                                                                                                                                                                                                                                                                                                  Zsebo KM, Suggs SV,
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                                                                                                                                                                                                                                                      (AMGE-) AMGEN INC.
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                                                           24-AUG-1990;
28-SEP-1990;
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Antibodies raised against the polypeptides are useful in diagnosis and to remove SCF from blood. This sequence represents SCF related DNA of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stem cell factor; SCF; haematopoietic activity; infertility; intestinal damage; myeloproliferative disorder; leucopenia; thrombocytopenia; anaemia; bone marrow transplant; immune deficiency; neoplasia; nerve damage; osteoporosis; metastatic carcinoma; leukaemia; miliary tuberculosis; haematopoietic progenitor cell; ss.
                                                                                                                                                                                         New non-natural stem cell factor, useful for treating e.g. leucopenia immune deficiency, also related nucleic acid and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                              Martin FH;
                                                                                                                                                                                                                            Disclosure; SEQ ID NO 32; 217pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stem cell factor (SCF) related DNA #33.
                                                                                                                                              Sugge SV,
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91US-00684535.
92US-00982255.
93US-00172329.
95US-00449653.
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                                                                                                                                             Zsebo KM, Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19; Conservative
                                                                                              BOSSELMAN R A.
                                                                                                                                                                  WPI; 2003-851459/79.
                                                                                                                     MART/) MARTIN F H.
                                                                                    ZSEBO K M.
                                                                                                           SUGGS S V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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01-OCT-1990;
10-APR-1991;
25-NOV-1992;
21-DEC-1993;
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                                                            12-JAN-1998;
                                                24-MAY-1995;
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                                                                                  ZSEB/)
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Matches
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The invention relates to stem cell factor (SCF) polypeptides with beematopoietic activity and the polynucleotides encoding them. The polypeptides are used for treating infertility, intestinal damage, myeloproliferative disorders, leucopenia, thrombocytopenia or anaemia, for improving engraftment of bone marrow transplants, for enhancing bone marrow tecovery after radiotherapy or chemotherapy and in treatment of immune deficiency, neoplasia, nerve damage, osteoporosis, metastatic carcinoma, leukaemia and miliary tuberculosis. The SCF polypeptides are also used to expand haematopoietic progenitor cells for transplantation and to prepare such cells for transfection with a gene. The SCF polymetic carcinoma and also as probes for mapping of the SCF gene, for identifying SCF-related diseases and as a marker for neighbouring genes. Antibodies raised against the polypeptides are useful in diagnosis and to remove SCF from blood. This sequence represents SCF related DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New non-natural stem cell factor, useful for treating e.g. leucopenia or immune deficiency, also related nucleic acid and antibodies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Suggs SV, Martin FH;
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                                                          90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
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90US-00537198.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Bosselman RA,
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Matches 19; Conservative
                                                                                                                                                                                                                                                                               ZSEBO K M.
BOSSELMAN R A.
                                                                                                                                                                                                                                                                               (ZSEB/) ZSEBO K M.
(BOSS/) BOSSELMAN R 1
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
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                                                                                             10-APR-1991;
25-NOV-1992;
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   11-JUN-1990;
24-AUG-1990;
                                                             01-OCT-1990;
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12-JAN-1998;
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31-OCT-2002

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, 5' or 3' end genomic flanking regions, 5' intrones of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, introcessive, and cytostatic activity. The composition may have a immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition may have a composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entiinflammatory steroid in a subject, for reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject stissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                       Aguilar D;
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                                                                                                                                                                                                                                                                                                                                           Pabalan J,
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                                                                                                                                                                                                                                                                                                                                           Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 554; 872pp; English.
                                                                                                                                                                                                                                                                                                                                       Nyce Jw, Li Y, Sandrasagra A, Ka
Miller S, Tang L, Shahabuddin S;
                                                                                23-APR-2002; 2002WO-US013135.
                                                                                                                                                                                                                                                           EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                     24-APR-2001; 2001US-0286137P
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Best Local Similarity 95.0°
Matches 19, Conservative
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31-OCT-2002
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ABZ89085;
                                                                                                                                                RESULT 942
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2708 TAAAAAAAAAAAAAAAAA 2727
                                                        20 TGAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                              Human oligonucleotide seguence.
                                                                                                                                                                                                              ABZ89301 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                             ABZ89301;
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RESULT 941

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ABZ89301

WO200285308-A2.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intradron codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention case in antisense gene therapy. The composition of the invention case in antisense gene therapy. The composition may have a comparating the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of continital ammatory steroid in a subject, for reducing levels of adenosine continital ammatory steroid in a subject, for reducing levels of adenosine comparation producing bronchodilation, increasing levels of adenosine contraction in subject is tissue, or treating bronchocomposition, clung surfactant in a subject, stissue, or treating bronchocomposition, clung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this parent is not represented in the printed continity in the subject of a respiratory disease or condition.
                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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Pred. No. 8.6e+02;
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                                                                                                                                                            Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 20 BP; 17 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 4543; 872pp; English.
                                                                                                                                                              Katz E,
                                                                                                                                                            Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
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                                    23-APR-2002; 2002WO-US013135.
                                                                              24-APR-2001; 2001US-0286137P.
                                                                                                                     (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95.0%;
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Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                      WPI; 2003-229219/22.
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                                                                                                                                                            Nyce JW, I
Miller S,
                                                                                                                                                                                                                                                                                                                                 ubiquinone
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24-APR-2001; 2001US-0286036P. 23-APR-2002; 2002WO-US013143

WO200285309-A2.

31-OCT-2002.

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding regaion, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory teroid and ubiquinone. A composition of the invention has antiinflammatory and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, ung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO.

"The sequence data for this patent is not represented in the printed specification, introducing format directly from WIPO.

"The sequence data for this patent is not represented in the printed specification, introducing commandation, in electronic format directly from WIPO.

"The sequence data for this patent is not represented in the printed specification, introducing producing producing and producing and producing and producing sequences are the printed specification, introducing sequences.
                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                          Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 4327; 872pp; English.
                                                                                                                                                                                                                          Katz E,
                                                                                                                                                                                                                 Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                     23-APR-2002; 2002WO-US013135.
                                                                                                           24-APR-2001; 2001US-0286137P.
                                                                                                                                                           (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                   WPI; 2003-229219/22.
  31-OCT-2002.
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                                                                                                                                                                                                                    Nyce Jw, I
Miller S,
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Sequence 20 BP; 19 A; 0 C; 1 G; 0 T; 0 U; 0 Other;

Gaps ö DB 1; Length 20; 1, Indels 0.7%; Score 18.4; DB 1; 95.0%; Pred. No. 8.6e+02; 0; Mismatches Local Similarity 95.0 nes 19; Conservative Query Match Matches

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## 2709 AAAAAAAAAAAAAAAA 2728 1 AGAAAAAAAAAAAAAAA 20

8 용 ABD25315 standard; DNA; 20 BP ABD25315; 

29-JUL-2004 (first entry)

AI092429-derived oligonucleotide SEQ ID 4327.

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfacent depletion; antiallergic, antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoonstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung alway or lung expression of a target polypeptide associated with lung alway or lung expression or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition composition has antiallergic, antialfammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lungs. The pulmonary obstruction, and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation, allergies, asthem, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the braskdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted truncleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                    Pabalan J, Aguilar D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                    Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 4327; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prevent any unwanted effects due to it
                                                                                                                                                                                                                                               Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                             bronchodilating agent.
                                                                                                                                                                                                                                                                                                                       WPI; 2003-093058/08.
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Miller S,
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2709 AAAAAAAAAAAAAAAA 2728 1 AGAAAAAAAAAAAAA 20 ABD21542 standard; DNA; 20 BP. (first entry) 19; Conservative 29-JUL-2004 ABD21542; RESULT 944 셤 XXXXXX ò

S100 calcium binding protein A2-derived oligo SEQ ID 554.

analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human, antisense, bronchoconstriction; allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity; lung, cancer, surfactant depletion, antiallergic, antiinflammatory, antiasthmatic;

Homo sapiens.

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; WPI; 2003-093058/08. Nyce JW, I Miller S,

Claim 15; SEQ ID NO 554; 763pp; English.

bronchodilating agent

comprising oligonucleotides, effective for alleviating occupance and one comparison describes and composition, alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, confident depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) adelivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adeling a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition as useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary veasconstriction, and or condition such as pulmonary disease, pulmonary condition replaces, asthma, impeded respiration, respiratory replaced to reduce the manner of the pulmorary inferriors and the manner of the pulmorary inferriors and the manner of the pulmorary inferriors and the pulmorary of the pu invention describes a novel composition (a) a first active agent, 

the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of

Sequence 20 BP; 1 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

ö Gaps ö 0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; Indels Pred. No. 8.6e; Mismatches ·; 95.0%; Conservative Best Local Similarity Matches 19; Conserv Query Match

20 TGAAAAAAAAAAAAAAA 1

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ABD2553 RESULT

ВР ABD25531 standard; DNA; 20

ABD25531;

29-JUL-2004 (first entry)

AI125651-derived oligonucleotide SEQ ID 4543.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory diserses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

Aguilar D;

Pabalan J,

Katz E,

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

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24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

ä Aguilar Pabalan J, Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-093058/08.

to Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4543; 763pp; English

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The followers of pulmonary obstruction, and/or lung and or available of the larget content or availability or malignant disease. In the lungs of the larget mRNA or to be pulmonary obstruction and/or lung and an entire lungs. inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of ngs.res

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06-MAY-2004
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                                                                                                                                                                                                                          Synthetic.
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                                                                             ADK67452;
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       RESULT 947
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence repersents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence the contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises an antisense oligonucleotides that are targeted
the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                          Human glucocorticoid receptor-specific antisense oligonucleotide #4234.
                                                                                                                                                  Gaps
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                                                                                                              DB 1; Length 20;
                                                                                                                                                1; Indels
                                                                      Sequence 20 BP; 17 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                        0.7%; Score 18.4; DB 1; ilarity 95.0%; Pred. No. 8.6e+02; Conservative 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 4234; 985pp; English.
                                                                                                                                                                                  2707 CTAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                    1 CTCAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                              ADH67400 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAY-2002; 2002US-0381857P.
                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                      Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
Les 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003099215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                ADH67400;
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Matches
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2708 TAAAAAAAAAAAAAAA 2727

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20 TCAAAAAAAAAAAAAAA 1

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The intercalator of the invention may be useful for the electrochemical detection of a gene, as an electrochemical double stranded DNA detection reagent and as an intercalator for inhibiting the influence of mismatch DNA and single stranded DNA. The intercalator enables the transmission of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel intercalator having a specific formula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            electronic transition between two base pairs to occur efficiently. The current sequence is that of the electrochemical detection intercalator related DNA 2 of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel intercalator, useful as electrochemical double stranded DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1776
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      intercalator; electrochemical detection; mismatch; ss.
                                                                                                                                     Electrochemical detection intercalator-related DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 19 A; 0 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 23; 24pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAAAA 2728
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ADK67452 standard; DNA; 20
                                                                                      (first entry)
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TUM KENKYUSHO KK.
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                                                                                                                                                                                                                                                                       JP2004024114-A.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navi.3, where the antisense compound a nucleic acid molecule encoding Navi.3, where the antisense compound appecifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathy, trigeminal neuropathy, arthritic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate pain for neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with human Navi.3 expression, the oligonucleotides are designed to target different regions of the human Navi.3 RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; omtidiabetic; microsomal prostaglandin E2 synthase inhibitor; cytoatatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; se.
                                                                                                       New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:654.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                           Claim 4; SEQ ID NO 1776; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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            14-AUG-2002; 2002US-0403416P
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*tag= b
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                                      (PHAA ) PHARMACIA CORP
                                                                                             WPI; 2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 19; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                  Roberds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 949
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to human mpGES-1 gate.

CC human mpGES-1 gene is located on chromosome 9, more specifically to having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human; antisense oligonucleotide; mitochondrial membrane; insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; ardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ophthalmic, immunological, cardiovascular or neurological disorder
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Pred. No. 8.6e+02;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ischaemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                               'note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 654; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     arthritis, diabetes, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2707 CTAAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2003; 2003WO-US030374.
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                                                                                                                 *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHAA ) PHARMACIA CORP
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Best Local Similarity
                                                                                                                                                                                                                                                         WO2004028458-A2
                                                                            modified base
                                                                                                                                                                                                                                                                                                                           08-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gierse JK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ischemia.
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useful for modulating

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The invention comprises antisense oligonucleotides that are targeted to the nucleic acids encoding a family of human proteins from mitochondrial membranes, which bind insulin sensitising, antidiabetic thiazolodinediones (referred to as mitoNEET). The antisense oligonucleotides of the invention are useful for modulating mitoNEET expression and for treating diseases or conditions associated with mitoNEET, such as diabetes, immunological disorders, cardiovascular disorders including hypertension, neurological disorders, and ischaemia/reperfusion injuries. The present DNA sequence represents a mitoNEET-specific antisense oligonucleotide of the invention. NOTE: The present sequence is a 2'-methoxyethyl (2'-MOE) gapmer with a
                                                                                                                                           New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aplastic anaemia, paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metaeratic carcinoma; acute leukaemia; myelosclerosis; osteopetrosis; metaeratic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Letterer-Siwe disease; coffectory erythroblastic anaemia; Di Guglielmo syndrome; congestive splenomegaly; Kala awar; sarcoidosis; primary splenic pancycopaenia; miliary tuberculosis; primary splenic pancycopaenia; miliary tuberculosis; vitamin Bl2 deficiency; folic acid deficiency; pyridoxine deficiency; vitaming blackfan anaemia; hypopigmentation disorder; piebaldism; vitiligo; neurological damage; infertility; intestinal damage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            irradiation; chemotherapy; AIDS; haematopoietic recovery; acute blood loss; neoplasm; cancer; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stem cell factor, SCF, universal PCR primer #4.
                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 87; 226pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 AAACAAAAAAAAAAAA 1
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90US-00537198.
90US-00573616.
90US-00589701.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-SEP-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 95.0
Marches 19; Conservative
(PHAA ) PHARMACIA CORP
                                                                                                 WPI; 2004-468836/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-OCT-1989;
11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JS6759215-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JUL-2004.
                                                 Colca JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP99304;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention comprises antisense oligonuclectides that are targeted to the nucleic acids encoding a family of human proteins from mitochondrial membranes, which bind insulin sensitising, antidiabetic thiazolodinadiones (referred to as: mitoNEET). The antisense oligonucleotides of the invention are useful for modulating mitoNEET expression and for treating diseases or conditions associated with mitoNEET, such as: diabetes, immunological disorders, cardiovascular disorders including hypertension, neurological disorders, and sischaemia/reperfusion injuries. The present DNA sequence represents a mitoNEET-specific antisense oligonucleotide of the invention. NOTE: The phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; antisense oligonucleotide; mitochondrial membrane; insulna sensitishing antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss;
                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mitoNEET-specific antisense oligonucleotide #87.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 1 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 141; 226pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2708 TAAAAAAAAAAAAAAA 2727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 TAAACAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADP69193 standard; DNA; 20 BP
                                                                                                                                                             25-NOV-2003; 2003WO-US037621.
                                                                                                                                                                                                          06-DEC-2002; 2002US-0431529P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-NOV-2003; 2003WO-US037621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-DEC-2002; 2002US-0431529P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-SEP-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 95.0°
Matches 19; Conservative
                                                                                                                                                                                                                                                         (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-468836/44.
                                                                WO2004053060-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2004053060-A2
                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUN-2004.
                                                                                                                24-JUN-2004
                                                                                                                                                                                                                                                                                                        Colca JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP69193;
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ADP69193,

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Gaps

Tue Nov

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The invention relates to preparing a (vertebrate) human stem cell factor (SCP) polyapetide comporting growing host cells transformed or transfected with DNA encoding a human SCF that simulates growth of transfected with DNA encoding a human SCF that simulates growth of transfected with DNA encoding a human SCF that simulates growth of polyapetide produced. Also included is a recombinant host cell cransformed or transfected with an expression control sequence, and isolating the polyapetide produced. Also included is a recombinant host cell cransformed or transfected with an expression control sequence, are retremented SCF polyapetide-encoding DNA operatively linked to a member of the vertebrate SCF polyapetide in the host cell. Instance of other nammals and recombinantly expressed SCF protein frame number of other mammals and recombinant techniques or for generating new and useful viral mannals and recombinant techniques or for generating new and useful viral cransferred prokaryotic and enkaryotic host cells, and new and useful transferred products. The DNA sequences are also useful as labelled probes in isolating human genomic DNA encoding SCF, in methods of probes in isolating human genomic DNA encoding SCF, in methods in developing transgenic mammalian species which may serve as enkaryotic of protein synthesis! In genetic therapy in humans and other mammals, and in developing transgenic mammalian species which may serve as enkaryotic notering proper in solating human genomic DNA encoding SCF, in methods in developing transgenic mammalian species which may serve as enkaryotic suseful for treating heameropoletic disorders, e.g., aplastic anammal, and encoding section of species which may serve as enkaryotic suseful for treating heameropoletic disorders, in developing transgenic and sections. The section of species which may serve as enkaryotic suseful for treating heameropoletic disorders, intenting promessing mannals, milary tuberculosis, milary tuberculosis, prinaman, and prypopigmentation or chemotherapy, an
                                                                                                                                                                                                                                                    Preparing a human stem cell factor (SCF) polypeptide, useful for treating
                                                                                                                                                                                                                                                                          hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                               Suggs SV, Martin FH;
                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 34; 210pp; English.
91US-00684535.
92US-00982255.
93US-00172329.
95US-00449182.
                                                                                                                                                               Bosselman RA,
                                                                                                                                                                                                          WPI; 2004-497128/47
                                                                                                                  (AMGE-) AMGEN INC.
                     25-NOV-1992;
21-DEC-1993;
24-MAY-1995;
  10-APR-1991;
                                                                                                                                                            Zsebo KM,
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ö Gaps ö 0.7%; Score 18.4; DB 1; Length 20; 55.0%; Pred. No. 8.6e+02; ve 0; Mismatches 1; Indels 95.0%; Conservative Best Local Similarity Matches 19; Conserv Query Match Matchea

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(first entry) 23-SEP-2004 ADP99302; XXXX

ADP99302 standard; DNA; 20

RESULT 953

ADP99302,

aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; multiple mystudy, moderner. Siwe disease;

refractory erythroblastic anaemia; Di Guglielmo syndrome;

congestive splenomegaly; Kala awar; sarcoidosis;

primary splenic pancytopaenia; miliary tuberculosis;

disseminated fungus disease; Fulminating septicaemia; malaria;

vitamin B12 deficiency; folic acid deficiency; pyridoxine deficiency;

biamond Blackfan anaemia; hypopigmentation disorder; piebaldism;

vitiligo; neurological damage; infertility; intestinal damage;

irradiation; chemotherapy; AIDS; haematopoietic recovery;

acute blood loss; neoplasm; cancer; ss; PCR; primer. cell; stem cell factor; gene therapy; haematopoietic progenitor cell factor, SCF, universal PCR primer #2 90US-00537198. 90US-00573616. 90US-00589701. 92US-00982255. 93US-00172329. 95US-00449182. 07-AUG-2000; 2000US-00635251. 91US-00684535. 89US-00422383 US6759215-B1 06-JUL-2004. 16-OCT-1989; 11-JUN-1990; 24-MAY-1995; 24-AUG-1990 10-APR-1991 25-NOV-1992 21-DEC-1993 Mammalia. 

Preparing a human stem cell factor (SCF) polypeptide, useful for treati hematopoletic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF. Suggs SV, Martin FH; Zsebo KM, Bosselman RA, WPI; 2004-497128/47.

(AMGE-) AMGEN INC.

Example 3; SEQ ID NO 32; 210pp; English.

The invention relates to preparing a (vertebrate) human stem cell factor (SCF) polypeptide comprising growing host cells transformed or transfected with DNA encoding a human SCF that stimulates growth of haematopoietic progenitor cells under nutrient conditions, the DNA being operatively linked to an expression control sequence, and isolating the operatively linked to an expression control sequence, and isolating the polypeptide produced. Also included is a recombinant host cell contransformed or transfected with an expression construct comprising a vertebrate SCF polypeptide encoding DNA operatively linked to a correbrate SCF polypeptide in the host cell. Disclosed as new are rat and human nucleic acids encoding SCF, SCF proteins from a number of other mammals and recombinantly expressed SCF proteins from a number of other mammals and recombinant techniques or for generating new and useful considering the large scale synthesis of SCF by a sequences are useful for effecting the large scale synthesis of SCF by a variety of recombinant techniques or for generating new and useful viral and circular plasmid DNA vectors. The DNA sequences are also useful as and circular plasmid DNA vectors. The DNA sequences are also useful as continued growth of such host cells capable of expression of the products of some products. The DNA sequences are also useful as confined probes in isolating human genomic DNA encoding SCF, in methods of production of SCF and SCF products in quantity. The SCF is the net of production of SCF and SCF products in quantity. The SCF is useful for treating hemaniana, myelofibrosis, myelogis, metastatic carcinoma, acute leukaemia, multiple myeloma, carcenoma, acute leukaemia, multiple myeloma, carcenoma, acute leukaemia, multiple myeloma, paccenome of production of such noar acute leukaemia, multiple myeloma, lymphoma, lymphoma, daucher's disease, lymphoma, daucher daucher daucher dauch

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888888888888888
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syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary splenic pancytopaenia, miliary tuberculosis, disseminated fungus disease, Fulminating septicaemia, malaria, vitamin B 12 and folic acid deficiency, pyridoxine deficiency, Diamond Blackfan annaemia, and hypopigmentation disorders such as piebaldism and vitiligo. The SCF are also useful for reating neurological damage, infertility states, intestinal damage resulting from irradiation or chemotherapy, and AIDS. SCF is also useful for enhancing haematopoietic recovery after acute blood loss and as a boost to the immune system for fighting neoplasia (cancer). The present sequence is a universal SCF primer used in the isolation of SCF DNA. Letterer-Siwe disease, refractory erythroblastic anaemia, Di Guglielmo

Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;

Score 18.4; DB 1; Length 20; Pred. No. 8.6e+02; 0; Mismatches 1; Indels 1; Indels 2707 CTAAAAAAAAAAAAAAAA 2726 20 CCAAAAAAAAAAAAAAA 1 Query Match Best Local Similarity 95.0%; 19; Conservative Matches ò ద

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RESULT 954

ADU50634 standard; DNA; 20 BP ADU50634; ADU50634/c 

13-JAN-2005 (first entry)

Human/rat stem cell factor, SCF, primer 220-11.

Stem cell factor; SCF; haematopoietic; HT1080 fibrosarcoma cell line; 5637 bladder carcinoma cell line; leukopaenia; thrombocytopaenia; anaemia; bone marrow during transplant; bone marrow aplasia; myeloguppression; immune deficiency; neoplasm; nerve damage; infertility; intestinal damage; myeloproliferative disorder; early haematopoietic progenitor cell; haematopoietic disorders; aplastic anaemia; myelofibrosis; myeloscierosis; osteopetrosis; metastatic carcinoma; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; biamond-Blackfan anaemia; DBA; Fanconi's anaemia; gene therapy; acute blood loss; ss; PCR; primer; probe.

Homo sapiens.

Rattus norvegicus.

US2004181044-A1.

16-SEP-2004.

19-JUN-2002; 2002US-00175608 16-OCT-1989

89US-00422383. 90US-00537198. 90US-00573616. 91US-00684535. 92US-009822255. 24-AUG-1990; 01-OCT-1990; 10-APR-1991; 11-JUN-1990

07-AUG-2000; 2000US-00635249 95US-00486546 21-DEC-1993; 25-NOV-1992;

BOSSELMAN R A. SUGGS S V. MARTIN F H. (ZSEB/) ZSEBO K M. (BOSS/) (MART/)

Martin FH; Suggs SV, Zsebo KM, Bosselman RA,

WPI; 2004-707481/69.

naturally occurring SCF, useful for treating leukopenia, thrombocytopenia, anemia, and enhancing engraftment of bone marrow during or stem cell factor (SCF) such as non-naturally-occurring SCF Novel

Example 3; SEQ ID NO 34; 216pp; English.

The invention relates to a stem cell factor (SCF) such as non-naturallycoccurring SCF having an amino acid sequence sufficiently duplicative of
that of naturally occurring SCF to allow possession of a hasanatopolatic
coccurring SCF haso included are an isolated DNA sequence for use in
securring SCF, Also included are an isolated DNA sequence for use in
coccurring SCF, Also included are an isolated DNA sequence for use in
securing SCF, Also included are an isolated DNA sequence cell of
transformed or transfected with the DNA, a polypeptide product of the
expression of the DNA in a prokaryotic or eukaryotic host cell
captession of the DNA in a prokaryotic or eukaryotic host
corrected DNA sequence coding for prokaryotic or eukaryotic host
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically functional plasmid or viral DNA vector
coll factor, a balologically planting stem cell factor, and an antibody (Ab)
corring the DNA, human SCF CDNA, obtained from HTJOBO
controms cell line, or human SCF CDNA, obtained from HTJOBO
controms cell line, or human SCF CDNA, obtained from propertions
coll factor, a baloning SCF. SCF is useful for treating patients
contivity of naturally occurring stem cell factor, and an antibody (Ab)
covortionam cell line, or human SCF CDNA, obtained from placed or covortion or myelosuppression which involves treating patients
contivity of naturally effective doses of SCF is useful for treating
coquired immune deficiency, neopleasi, or chemotherapeutic for functional patients
companies of the cultured cells with a gene SCF is useful for
cransfecting a gen useful for treating various haematopoietic disorders, appastic anaemia, myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, Niemann-Pick disease, Diamond-Blackfan anaemia (DBA), Fanconi's anaemia. SCF is useful for enhancing the efficiency of gene therapy, for enhancing haematopoietic recovery after acute blood loss. The present sequence is a primer and/or probe used in the isolation of SCF nucleic

Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

ö 'Match 0.7%; Score 18.4; DB 1; Length 20; Local Similarity 95.0%; Pred. No. 8.6e+02; les 19; Conservative 0; Mismatches 1; Indels Query Match Matches

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2707 CTAAAAAAAAAAAAAA 2726 20 CGAAAAAAAAAAAAAAA 1 ADU50632 standard; DNA; 20 BP. RESULT 955 ADU50632/c 셤 ò

Human/rat stem cell factor, SCF, primer 220-3. (first entry) 13-JAN-2005 

ADU50632;

Stem cell factor; SCF; haematopoietic; HT1080 fibrosarcoma cell line; 5637 bladder carcinoma cell line; leukopaenia; thrombocytopaenia;

Novel stem cell factor (SCF) such as non-naturally-occurring SCF or naturally occurring SCF, useful for treating leukopenia, thrombocytopenia, anemia, and enhancing engraftment of bone marrow during anaemia; bone marrow during transplant; bone marrow aplasia; myelosuppression; immune deficiency; neoplasm; nerve damage; infertility; intestinal damage; myeloproliferative disorder; early haematopoletic progenitor cell; haematopoletic disorders; aplastic anaemia; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Diamond-Blackfan anaemia; DBA; Fanconi's anaemia; gene therapy; acute blood loss; ss; PCR; primer; Martin FH; Sugge SV, 89US-00422383. 90US-00537198. 90US-00573616. 90US-00589701. 91US-00684535. 93US-00172329 95US-00486546 2000US-00635249 19-JUN-2002; 2002US-00175608 Bosselman RA, ZSEBO K M. BOSSELMAN R A. WPI; 2004-707481/69. MARTIN F H. Rattus norvegicus. JS2004181044-A1. transplantation. SUGGS S 07-AUG-2000; Homo sapiens 21-DEC-1993; 07-JUN-1995; 10-APR-1991, 25-NOV-1992, 16-OCT-1989 24-AUG-1990 16-SEP-2004 11-JUN-1990 01-OCT-1990 Zsebo KM, (SUGG/) (MART/) ZSEB/) (Boss/)

Example 3; SEQ ID NO 32; 216pp; English.

The invention relates to a stem cell factor (SCF) such as non-naturallycocurring SCF having an amino acid sequence sufficiently duplicative of
that of naturally occurring SCF to allow possession of a haemacopoietic
biological activity of naturally occurring stem cell factor, or naturally
cocurring SCF. Also included are an isolated DNA sequence for use in
securing expression in a prokaryotic or eukaryotic host cell of nontransformed or transfected with the DNA, a polypeptide product of the
expression of the DNA in a prokaryotic or eukaryotic host
corrected DNA sequence coding for prokaryotic or eukaryotic host
cell factor, a biologically functional plasmid or viral DNA vector
cell factor, a biologically functional plasmid or viral DNA vector
cell factor, a biologically functional plasmid or viral DNA vector
cell factor, a biologically functional plasmid or viral DNA vector
cell factor, a biologically functional plasmid or viral DNA vector
coll factor, a biologically functional plasmid or viral DNA vector
cell factor, a biologically functional plasmid or viral DNA vector
coll factor, a biologically functional plasmid or viral DNA vector
coll factor, a biologically functional plasmid or viral DNA vector
coll factor, a biologically cannown or coll ince and plasmid or viral DNA, a polypeptide having part
colluding the DNA sequence abcoded by composite multiple of a stably transformed or transfected with the DNA, a polypeptide having part
corrections a cell line, or human SCF cDNA obtained from H71080
corrections cell factor, and an antibody (Ab)
specifically binding SCF. SCF is useful for treating leukopaenia,
cactivity of naturally-occurring stem cell factor, and an antibody (Ab)
specifically binding SCF. SCF is useful for treating plantating patients
covery in treatment of radiation, chemical, or chemotherapeutic induced
covery in treatment of radiation, chemical, or chemotherapeutic or may be sequence
covery in treatment of factive does of SCF is useful for treating
acquired immune deficiency, neoplasia, ner

ö transfecting early haematopoietic progenitor cells with a gene which involves culturing early haematopoietic progenitor cells with SCF, and transfecting the cultured cells with a gene. SCF is useful for transfecting a gene to a mammal which involves culturing early haematopoietic progenitor cells with SCF, transfecting the cultured cells with a gene, and administering the cultured cell to the mammal. SCF is useful for treating various haematopoietic disorders, aplastic anaemia, myelofibrosis, myelosclerosis, osteopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, Hodgkin's disease, lymphoma, Gaucher's disease, Niemann-Pick disease, Diamond-Blackfan anaemia (BBA), Fanconi's anaemia. SCF is useful for enhancing the efficiency of gene therapy, for enhancing haematopoietic recovery after acute blood loss. The present sequence is a primer and/or probe used in the isolation of SCF nucleic Stimulating proliferation of melanocyte cells in human, involves administering stem cell factor polypeptide or its biologically active fragments stimulating growth of melanocyte cells, and optionally carrier, to human. Antianemic; Antiemetic; Cytostatic; Anti-HIV; Cardiovascular-Gen.; CNS-Gen.; Antibarasitic; Antibacterial; Immunosuppressive; Antiharinflammatory; Fungicide; Antifertility; AIDS; aplastic anemia; parcoxysmal nocturnal hemoglobinuria; osteopetrosis; acute leukemia; multiple myeloma; hodgkins disease; lymphoma; gauchers disease; niemann pick disease; sarcoidosis; plasmodium infection; vitamin deficiency; hypopigmentation; vitiligo; infertility; chronic myelocytic leukemia; cell proliferation; Stem Cell Factor; PCR; primer; ss. Gaps ö Length 20; 1; Indels Universal Stem Cell Factor PCR primer 220-3, SEQ ID 32. Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other; Score 18.4; DB 1; Pred. No. 8.6e+02; Bosselman RA, Suggs SV, Martin FH; 0; Mismatches 2707 CTAAAAAAAAAAAAAAA 2726 20 CCAAAAAAAAAAAAAAAA 1 89US-00422383. 90US-00537198. 90US-00573616. 90US-00589701. 91US-00684535. 92US-00982255. 93US-00172329. 95US-00449649. BP. 0.7%; 2000US-00604325 ADW93077 standard; DNA; 20 (first entry) Local Similarity 95.0 WPI; 2005-160562/17. (AMGE-) AMGEN INC. US6852313-B1. 26-JUN-2000; 10-APR-1991; 25-NOV-1992; 21-DEC-1993; 21-APR-2005 16-OCT-1989; 38-FEB-2005 11-JUN-1990 24-AUG-1990 01-OCT-1990 24-MAY-1995 Zsebo KM, Synthetic. ADW93077; Query Match acids. Matches ADW93077, RESULT ð 셤

89US-00422383. 90US-00537198. 90US-00573616.

Example 3; SEQ ID NO 32; 212pp; English.

7 10:41:34 2006

Tue Nov

90US-00589701. 91US-00684535. 92US-00982255. 93US-00172329.

95US-00449649

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Antianemic; Antiemetic; Cytostatic; Anti-HIV; Cardiovascular-Gen.; CNS-Gen.; Antiparastitic; Antibacterial; Immunosuppressive; Antihacterial; Immunosuppressive; Antiinflammatory; Fungicide; Antifertility; AIDS; aplastic anemia; paroxysmal nocturnal hemoglobinuria; osteopetrosis; acute leukemia; multiple myeloma; hodgkins disease; lymphoma; gauchers disease; niemann pick disease; sarcoidosis; plasemodium infection; vitamin deficiency; hypopigmentation; vitiligo; infertility; chronic myelocytic leukemia; cell proliferation; Stem Cell Factor; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                 20 CCAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                              2707 CTAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                       .079/c
ADW93079 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                      invention for cloning SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-APR-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6852313-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          38-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADW93079;
                                                                                                                                                                                                                                                                                              therapy
                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 957
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Stimulating proliferation of melanocyte cells in human, involves administering stem cell factor polypeptide or its biologically active fragments stimulating growth of melanocyte cells, and optionally carrier,
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                                                                                                                                                                                                                                                                                Zsebo KM, Bosselman RA,
                                                                                                                                                                                                                                                                                                                                  WPI; 2005-160562/17.
                                                                                                                                                                                                                                (AMGE-) AMGEN INC.
                                                                                                                                                   21-DEC-1993;
24-MAY-1995;
                          11-JUN-1990;
24-AUG-1990;
                                                                         01-OCT-1990;
                                                                                                       10-APR-1991
                                                                                                                            25-NOV-1992
The present invention relates to a method (M1) for stimulating proliferation of melanocyte cells in a human. (M1) involves administering a Stem Cell Factor (SCF) protein, or its biologically active fragments that stimulates growth of melanocyte cells, and optionally a carrier, to the human. The SCF is covalently conjugated to a water soluble polymer of the human. The SCF is covalently conjugated to a water soluble polymer of the human. The SCF is covalently conjugated to a water soluble polymer of more other cytckines. SCF is also able to stimulate the growth of primitive progenitors such as early hematopoietic progenitor cells that care capable of maturing to erythroid, megakaryccyte, granulocyte, configurating to erythroid, megakaryccyte, granulocyte, configurating bone marrow regeneration, and in augmenting T cells neural stem cells and primordial germ stem cells (M1) is useful in treating AIDS, configurating bone marrow regeneration, and in augmenting T cell production. (M1) is useful in treating AIDS, aplastic anemia, parcoxysmal nocturnal hemoglobinuria, myelofibrosis, aplastic anemia, parcoxysmal nocturnal hemoglobinuria, cute leukemia, myelostaronia, disseminated fungus disease, fulminating myelosma, Hodgkin's disease, lymphoma, Gaucher's disease, congestive splenomegaly, Kalaacar, sarcoidosis, primary splenic pancytopenia, disseminated fungus disease, fulminating septicemia, malaria, vitamin B12 and folic acid deficiency disease, pridebaldism and vitiligo. (M1) is useful in treating infertility states, intestinal damage resulting from irradiation or chemotherapy, and stem cell myebbaldism and vitiligo. (M1) is useful in treating infertility states, primary thrombocythemia and acute leukemia. (M1) is useful in syngeneic or autologous contents and acute leukemia. (M1) is useful in syngeneic or autologous from primary thrombocythemia and acute leukemia. (M1) is useful in an example from in equipance, is an eleukemia in an example from in enemental marow tangeneic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.7%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 8.6e+02; Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Martin FH;

Suggs SV,

The present invention relates to a method (M1) for stimulating proliferation of melanocyte cells in a human. (M1) involves administering composition of melanocyte cells in a human. (M1) involves administering to a Stem Cell Factor (SCF) protectain, or its biologically active fragments that stimulates growth of melanocyte cells, and optionally a carrier, to the human. The SCF is covaligated to a water soluble polymer e.g. polyethylene glycol. Also, the SCF is co-administered with one or covare other cytokines. SCF is also able to stimulate the growth of primitive progenitors such as early hematopoietic progenitor cells that are consequent of maturing to erythroid, megakaryocyte, granulocyte, composition of maturing to erythroid, megakaryocyte, granulocyte, composition macrophage cells, and non-hematopoietic stem cells such as neutal stem cells and primordial germ stem cells. (M1) is useful in accelerating bone marrow regeneration, and in augmenting T cell accelerating bone marrow regeneration, and in augmenting T cell accelerating bone marrow regeneration, and in augmenting T cell accelerating bone marrow regeneration in functional marrow mass due to toxic, radiant or immunological injury. (M1) is useful in treating AlDS, conjected by a reduction in functional marrow mass due to toxic, radiant or immunological injury. (M1) is useful in treating AlDS, conjecters, osteoperosis, metastatic carcinoma, acute leukemia, mylosic mylosic carcinoma, acute leukemia, conjecters on orgestive splenomegaly, Kalazar, sarcoidosis, primary cylosic pancycopenia, disease, lymphoma cate deficiency disease, and hypopigmentation disorders such as choric myelogenes, conjective pancycopenia, disease, and hypopigmentation deficiency disorders such as choric myelogenia, thrombocythemia and acute leukemia. (M1) is useful in treating infertility states, intestinal damage resulting from irradiation or chemotherapy, and progenitors in syngeneic, allogeneic or autologous cell myelomia and autore leukemia. (M1) is useful in an example from irra Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other; therapy. The present seque invention for cloning SCF. Universal Stem Cell Factor PCR primer 220-11, SEQ ID 34. 2726

Gaps ò 0.7%; Score 18.4; DB 1; Length 20; 15.0%; Pred. No. 8.6e+02; ve 0; Mismatches 1; Indels 2707 CTAAAAAAAAAAAAAAAA 2726 20 CGAAAAAAAAAAAAAAA 1 ADZ47531 standard; DNA; 20 BP. 92.08; 19; Conservative Query Match Best Local Similarity RESULT 958 Matches ADZ47531/ 셤 ઠ

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Universal PCR primer, 220-11, SEQ ID NO: 34. 30-JUN-2005 (first entry)

ADZ47531;

EXEXEXE Executive

26-JUN-2000; 2000US-00604325.

Stem cell factor; cell growth; immune disorder; immunomodulator; genetic disorder; hematological disease; antianemic; cancer; cytostatic; enceptasm; neurological disease; neuroprotective; infection; antimicrobial; hypopigmentation; dermatological; infertility; antifinertility; inflammation; antiinfertility; gene therapy; PCR;

Universal PCR primer, 220-3, SEQ ID NO: 32.

(first entry)

30-JUN-2005

ADZ47529;

ADZ47529 standard; DNA; 20 BP.

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RESULT 959
             ADZ47529/c
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The present invention relates to a method of stimulating growth of stromal cells in a human. The method involves administering to the human at complete and an effective amount of a human stem cell factor (SCP) polypeptide and optionally a pharmaceutically acceptable carrier. The invention is useful for treating immune disorders such as (acquired immune deficiency as niemann pick disease), hematological diseases (such as multiple myeloma, hodgkins disease), neurological diseases (such as multiple myeloma, infection (such as leishmania donovani infection), hypopigmentation disorders (such as plebaldism and vitiligo), in the treatment of infection or chemotherapy and inflammation (such as sarcoidosis). The invention is also useful in gene therapy and SCP therapy. The present control of the an universal PCR primer. This primer is used in the cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stimulating growth of stromal cells for treating AIDS or severe combined immunodeficiency states in human, primary splenic pancytopenia, miliary tuberculosis, by administering human stem cell factor polypeptide and carrier to human.
Stem cell factor; cell growth; immune disorder; immunomodulator; genetic disorder; hematological disease; antianemic; cancer; cytostatic; neoplasm; neurological disease; neuroprotective; infection; antimicrobial; hypopigmentation; dermatological; infertility; antinfertility; lnflammation; antiinflammatory; gene therapy; PCR;
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0.7%; Score 18.4; DB 1; Length 20;
Best Local Similarity 95.0%; Pred. No. 8.6e+02;
Matches 19; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 34; 216pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  900S-00537198
900S-00573616.
900S-00589701.
910S-00684535.
920S-00172325.
930S-00172329.
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                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUL-2003; 2003US-00620642
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BOSSELMAN R A.
SUGGS S V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2005-321855/33.
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                                                                                                                                                                                                                                                                                                  US2005080250-A1.
                                                                                                                                                                                                                                   Unidentified
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21-DEC-1993;
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10-APR-1991;
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                                                                                                                                                                          primer; ss.
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(SUGG/)
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90US-00537198. 90US-00573616. 90US-00589701. 91US-00684535. 92US-00982255.

10-APR-1991; 25-NOV-1992;

01-OCT-1990, 24-AUG-1990

21-DEC-1993; 07-JUN-1995;

16-JUL-2003; 2003US-00620642

US2005080250-A1. Unidentified.

primer; ss.

14-APR-2005

07-JUN-1995; 95US-00486546. 07-AUG-2000; 2000US-00635249. 19-JUN-2002; 2002US-00175608.

Suggs SV, Martin FH;

Bosselman RA,

Zsebo KM,

WPI; 2005-321855/33.

BOSSELMAN R A. SUGGS S V.

BOSS/)

(SUGG/) SUGGS S V. (MART/) MARTIN F H. (ZSEB/) ZSEBO K M.

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The present invention relates to a method of stimulating growth of stromal cells in a human. The method involves administering to the human stromal cells in a human. The method involves administering to the human confidencive amount of a human stem cell factor (SCF) polypeptide and optionally a pharmaceutically acceptable carrier. The invention is useful for treating immune disorders such as (acquired immune deficiency syndrome, severe combined immunodeficiency), genetic disorder (such as niemann pick disease), hedisease, anemia), cancers (such as multiple myeloma, hodgkins disease, spleen disease, anemia), cancers (such as acute infection (such as leishmania donovani infection), hypopigmentation disorders (such as plebaldism and vitiligo), in the treatment of infertility states and for treating intestinal damage resulting from irradiation or chemotherapy and inflammation (such as sarcoidosis). The invention is also useful in gene therapy and SCF therapy. The present sequence is an universal PCR primer. This primer is used in the cloning
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stimulating growth of stromal cells for treating AIDS or severe combined immunodeficiency states in human, primary splenic pancytopenia, miliary tuberculosis, by administering human stem cell factor polypeptide and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; SEQ ID NO 32; 216pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rat and human SCF cDNA.
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nes 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       carrier to human.
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Gaps

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The present invention relates to a method (M1) for labeling double stranded nucleic acid for efficient detection of DNA or RNA. The method comprises using a carbodismide derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid. (M1) is useful for labeling double stranded or single stranded nucleic acid or detecting single nucleotide polymorphisms. The present sequence was used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                     Labeling double stranded nucleic acid, involves utilizing carbodiimide derivative for labeling thymine, uracil and guanine, which exists in mismatch region of nucleic acid or unstable region of hydrogen bond of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibody; stem cell factor; probe; PCR; primer; ss.
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                                                                                                                                                                                                                    Tabata E;
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900S-00537198
900S-0059701
910S-00684535
920S-00982255
930S-0048729
2000US-00643659.
                                                                                                           18-MAR-2004; 2004JP-00078900
                                                                                                                                              18-MAR-2004; 2004JP-00078900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEE01384 standard; DNA; 20
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                                                                                                                                                                                                                 Fakenaka S, Nojima
                                                                                                                                                                                                                                                    WPI; 2005-685344/71
                                                                                                                                                                               (TAKE/) TAKENAKA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                      JP2005265617-A.
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25-NOV-1992;
21-DEC-1993;
                                                                                                                                                                                                                                                                                                                                              nucleic acid
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21-AUG-2000;
                                                                          29-SEP-2005.
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24-AUG-1990;
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     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method (M1) for labeling double stranded nucleic acid for efficient detection of DNA or RNA. The method comprises using a carbodininde derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid or defecting labeling double stranded or single stranded nucleic acid or detecting single nucleotide polymorphisms. The present sequence was used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Labeling double stranded nucleic acid, involves utilizing carbodiimide derivative for labeling thymine, uracil and guanine, which exists in mismatch region of nucleic acid or unstable region of hydrogen bond of
                                                                                                                                                                                                                              Oligonucleotide ODN2 used to illustrate nucleic acid labeling method
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                                                                                                                                                                                                                                                                 DNA detection; RNA detection; SNP detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 18.4; DB 1; 95.0%; Pred. No. 8.6e+02;
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CTAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                            (first entry)
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nes 19; Conservative
                                                                                                                    AED13294 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                     JP2005265617-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
                                                                                                                                                                                                                                                                                                    Synthetic
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Query Match

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Gaps

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1; Indels

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AEE60694,
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                                                                                                                                                                                                                     The invention relates to a purified antibody that is specifically immunoreactive with a stem cell factor (SCF) or SCF receptor. Also described: (1) a hybridoma cell line producing a monoclonal antibody that is specifically immunoreactive with a SCF protein; (2) inhibiting the activity of a mast cell population; (3) decreasing blood cell proliferation, maturation or activity in a in a mammal; (4) decreasing the interaction between a SCF and an SCF receptor in a cell population; (5) treating a mammal having a disorder mediated through the interaction of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF polypeptide, and a pharmaceutically immunoreactive with an SCF comprised and methods are useful for inhibiting the activity of a mast cell population, decreasing the interaction maturation or activity in a in a mammal, decreasing the interaction between a SCF and an SCF receptor in a cell population, and treating a mammal having a clivity of maturation or activity in a in a mammal, decreasing the interaction of SCF with an SCF receptor. The antibody, composition, and methods are useful for treating a mammal having a clistoder mediated through the interaction of SCF with an SCF receptor. The antibody, composition, and methods are useful for treating disorders, cuberculosis, or malaria. The present sequence represents a universal cligonucleotide sequence which can be used as a probe or a PCR primer in the amplification and sequencing of rat and human SCF, which is used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                          New stem cell factor antibody, useful for treating hematopoietic disorders such as anemia, leukemia, lymphoma, HIV, tuberculosis, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.7%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 8.6e+02; Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibody; stem cell factor; probe; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                  Martin FH;
                                                                                                                                                                                                  Example 3; SEQ ID NO 34; 217pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Universal oligonucleotide SEQ ID NO:32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 example from the present invention.
                                                                                  Sugge SV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2707 CTAAAAAAAAAAAAAAA 2726
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90US-00537198.
90US-00573616.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEE01382 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                    Bosselman RA,
              ZSEBO K M.
BOSSELMAN R A.
              (ZSEB/) ZSEBO K M.
(BOSS/) BOSSELMAN R ?
(SUGG/) SUGGS S V.
(MART/) MARTIN F H.
                                                                                                              WPI; 2005-796179/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2005261175-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-1989;
11-JUN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JAN-2006
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                                                                                    Zsebo KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEE01382;
                                                                                                                                                                         malaria.
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The invention relates to a purified antibody that is specifically immunoreactive with a stem cell factor (SCF) or SCF receptor. Also described: (1) a hybridoma cell line producing a monoclonal antibody that is specifically immunoreactive with a SCF protein; (2) inhibiting the activity of a mast cell population; (3) decreasing blood cell proliferation, maturation or activity in a in a mammal; (4) decreasing the interaction between a SCF and an SCF receptor in a cell population; (5) treating a mammal having a disorder mediated through the interaction of SCF with an SCF receptor; and (6) a pharmaceutical composition of SCF with an SCF receptor; and (6) a pharmaceutical composition comprising an antibody specifically immunoreactive with an SCF antibody and methods are useful for inhibiting the activity of a mast cell population, decreasing blood cell proliferation, maturation or activity in a in a mammal, decreasing the interaction between a SCF and SCF receptor in a cell population, and treating a mammal having a disorder mediated through the interaction of SCF with an SCF receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The antibody, composition, and methods are useful for treating disorders, e.g. hematopoietic disorders such as anemia, leukemia, lymphoma, HIV, tuberculosis, or malaria. The present sequence represents a universal oligonucleotide sequence which can be used as a probe or a PCR primer in the amplification and sequencing of rat and human SCF, which is used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New stem cell factor antibody, useful for treating hematopoietic disorders such as anemia, leukemia, lymphoma, HIV, tuberculosis, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                              Suggs SV, Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 32; 217pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        an example from the present invention
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                           92US-00982255.
93US-00172329.
95US-00448729.
91US-00684535
                                                                                                                               21-AUG-2000; 2000US-00643659
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                                                                                                                                                                                                                                                                                                                                                                  Bosselman RA,
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Best Local Similarity
                                                                                                                                                                                                ZSEBO K M.
BOSSELMAN R
                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2005-796179/81.
                                                                                                                                                                                                                                                                                                    (MART/) MARTIN F H.
                                                                                                                                                                                                                                                                  SUGGS S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6967029-B1.
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10-APR-1991;
25-NOV-1992;
                                                          21-DEC-1993;
24-MAY-1995;
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                                                                                                                                                                                                                                                                                                                                                                     Zsebo KM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      malaria.
                                                                                                                                                                                                    (ZSEB/)
                                                                                                                                                                                                                                                                  (snee/)
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The invention relates to a method (M1) for enhancing hematopoiesis in a human or other subject. (M1) comprises: (a) obtaining hematopoietic content subject. (M1) comprises: (a) obtaining hematopoietic content step (a) by adding to the call a stem cell factor (SCF) obtained in step (a) by adding to the call a stem cell factor (SCF) cobtained in step (a) by adding to the call a stem cell factor (SCF) cobtained in step (b) therefore restoring fragments that stimulate growth of hematopoietic progenitor cells; and (c) administering to the human or other subject hematopoietic progenitor cells and enhancing content subject in the human or other subject and enhancing contents in the human or other subject and enhancing comprises: (a) obtained hematopoietic progenitor cells a method (M2) for expanding hematopoietic progenitor cells at a method comprises: (a) obtained in step (b) therefore restoring hematopoietic progenitor cells as well from a donor; and (b) expanding the cells obtained in step (a) by adding to the cells the comprises: (a) obtained in step (a) by adding to the cells the content subject. (M2) is useful for expanding hematopoietic progenitor cells, where the hematopoietic cells are chosen from stem cells of progenitor cells, where the hematopoietic cells are chosen from stem cell deficiencies such as aplastic anemia, paroxymal coturnal hemoglobinuria, myelofibrosis, myelosclerosis, Gaucher's cilsease, Nuleman-Pick disease, Hodgkin's disease, Kala-azar, sarcoidosis, disease, Mu2) is useful in expanding early hematopoietic progenitors. The conemia, hypopigmentation disorders such as piebaldism and vitiligo, and Albs., and folic acid deficiency, pyridoxine deficiency, piamond blacken squence represente a universal PCR primer for SCPP, which is used in an example from the event in an evament of the event in a universal PCR primer for sequence represente a universal p
                                                                                                                                                                                                                                                                                                                                                                                                                      Enhancing hematopoiesis in human, comprises expanding hematopoietic progenitor cells by adding stem cell factor polypeptide to cells and administering expanded cells to human.
                                                                                                                                                                                                                                                                                                                               Martin FH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; SEQ ID NO 32; 213pp; English.
                                                                                                                                                                                                                                                                                                                               Suggs SV,
                                                                                       90US-00537198.
90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
                        21-AUG-2000; 2000US-00643659.
                                                                                                                                                                                                                                                                                                                             Bosselman RA,
                                                                                                                                                                                                                                                                                                                                                                           WPI; 2006-053612/06.
                                                                                                                                                                                                                                                                               (AMGE-) AMGEN INC.
                                                                                                                                                          10-APR-1991;
25-NOV-1992;
21-DEC-1993;
                                                                                       11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
                                                                                                                                                                                                                                 24-MAY-1995;
                                                                 16-OCT-1989
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Sequence 20 BP; 0 A; 0 C; 2 G; 18 T; 0 U; 0 Other;

Gaps ö Query Match 0.7%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 8.6e+02; Matches 19; Conservative 0; Mismatches 1; Indels

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AEE60696 standard; DNA; 20 BP AEE60696; RESULT 965 **AEE60696**,

(first entry)

09-FEB-2006

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Universal stem cell factor PCR primer SEQ ID NO:34.
             hematopoiesis; stem cell factor; PCR; primer; ss.
                                                                                                            Bosselman RA, Suggs SV, Martin FH;
                                                                       90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
                                                              90US-00537198.
                                                 2000US-00643659
                                                          89US-00422383
                                                                                          95US-00449649
                                                                                                    (AMGE-) AMGEN INC.
                               US6967029-B1
                                                 21-AUG-2000;
                                        22-NOV-2005.
                                                                                          24-MAY-1995;
                                                          16-0CT-1989;
                                                               11-JUN-1990;
24-AUG-1990;
                                                                                25-NOV-1992
                                                                        01-OCT-1990
                                                                             10-APR-1991
                      Synthetic.
                                                                                                            Zsebo KM,
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Enhancing hematopoiesis in human, comprises expanding hematopoietic progenitor cells by adding stem cell factor polypeptide to cells and administering expanded cells to human. WPI; 2006-053612/06.

Example 3; SEQ ID NO 34; 213pp; English.

The invention relates to a method (M1) for enhancing hematopoiesis in a human or other subject. (M1) comprises: (a) obtaining hematopoietic continuant or other subject; (b) expanding the cells obtained in step (a) by adding to the cell a stem cell factor (SCF) cobtained in step (a) by adding to the cell a stem cell factor (SCF) cobtained in step (a) by adding to the cell a stem cell factor (SCF) coptained in step (b) to respanded hematopoietic progenitor cells a ministering to the human or other subject the expanded hematopoietic progenitor cells continuated in step (b). Therefore restoring hematopoietis to effect comprises in the human or other subject. Also described is a method (M2) for expanding hematopoietic progenitor cells a method (CC comprises: (a) obtaining hematopoietic progenitor cells a method (CC comprises: (a) obtaining hematopoietic progenitor cells witch a donor; and (b) expanding the cells obtained in step (a) by adding to the cells the SCF polypeptide or its biological active fragments. (M1) is useful for expanding hematopoietic progenitor cells, megakaryocytes and erythroblates. (M2) is useful for expanding hematopoietic progenitor cells, megakaryocytes and erythroblates. (M1) is useful for cells, megakaryocytes and erythroblates. (M1) is useful for treating various stem cell deficiencies such as aplastic anemia, paroxymal cc disseminated fungus disease, Hodghir's disease, Kala-azar, sarcoidosis, disease, Niemann-Pick disease, Hodghir's disease, Mala-azar, sarcoidosis, CC disseminated fungus disease, Hodghir's disease, with an alaria, vitamin Black (M2) is useful in expanding early hematopoietic progenitors in summarces hematopoietic and servanding candedicion disorders such as piebaldism and vitiligo, and canemia, hypopigmentation disorders such as piebaldism and vitiligo, enhances hematopoietic progenitors enhances hematopoietic progenitors enhances hematopoietic proderic propenitor enhances hematopoietic proderic proderic proderic proderic proderic proderic proderic proderic proderic prod enhances hematopolesis by expanding early hematopoletic progenitors. The present sequence represents a universal PCR primer for SCF, which is used in an example from the present invention.

Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 8.6e+02; ive 0; Mismatches 1; Indels Local Similarity 95.0 nes 19; Conservative Query Match Best Loca Matches

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2707 CTAAAAAAAAAAAAAAA 2726

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AEF16083

RESULT 966 AEF16083,

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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                 gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded CDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     yene expression; reverse transcription; primer; cDNA; restriction enzyme; 89.
                               Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 2 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
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Pred. No. 8.8e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                         by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 8; 11pp; Japanese.
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Best Local Similarity 95.03
Matches 19; Conservative
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                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a novel isolated nucleic acid molecule capable of modulating transcription, where the nucleic acid molecule shows at least 80 % sequence identity to a sequence listed in Table 1 of the specification (or its complement). The nucleic acid molecule is useful for identifying promoters comproments, or fragments to modulate transcript levels in plants. The present sequence is that of a plant promoter DNA sequence of the invention (derived from either Arabidopsis thaliana or Oryza sativa, but not clearly stated in the specification) which is claimed as being capable of modulating transcription.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcription,
                                                                                                                                                                                                 Transcription modulation-related plant promoter DNA sequence SeqID71.
                                                                                                                                                                                                                                  DNA purification; gene expression; crop improvement;
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useful for identifying promoters, control elements, or fragments
modulate transcript levels in plants.
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OKAMURO J.
SCHNEEBERGER F
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KWOK S.
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                  Disclosure; Page 6; 11pp; Japanese.
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AAQ75676 standard; DNA; 21
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Disclosure; Page 9; 11pp; Japanese
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Matches 19; Conservative
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Pred. No. 8.8e+02;
0; Mismatches 1; Indels
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 8.8e+02;
0; Mismatches 1; Indels
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                            AAQ75650 standard; DNA; 21 BP.
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by digestion with restriction enzymes.
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Pred. No. 8.8e+02;
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BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
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                                   / Match 0.7%; Score 18.4; DB 1; Local Similarity 95.0%; Pred. No. 8.8e+02; nes 19; Conservative 0; Mismatches 1;
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         A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded CDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of CDNAs, in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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AAQ75775 standard; DNA; 21
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Best Local Similarity 95.0
Matches 19; Conservative
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AAQ75697/
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Pred. No. 8.8e+02;
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Best Local Similarity 95.0
Matches 19; Conservative
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Best Local Similarity
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Pred. No. 8.8e+02;
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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95.0%; Pred. No. 8.8e+02;
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(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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A method for the analysis of cDNA comprises (a) preparing an aggregate of
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AAQ75774/c
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  Score 18.4; DB 1;
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Length 21; 1; Indels

Score 18.4; DB 1; Pred. No. 8.8e+02; 0; Mismatches

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- by amplification of mRNA followed

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expression; reverse transcription; primer; cDNA;

restriction enzyme; ss.

(first entry)

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Reverse transcription primer used in cDNA analysis technique.
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          AAQ75745 standard; DNA; 21 BP.
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Pred. No. 8.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ75677;
                                                                                                                                                                                                  Query Match
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of
Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                        A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis; gene expression; reverse transcription; primer; cDNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                             0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02;
                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                       Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                          Disclosure, Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                         2706 ACTAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                               ВЪ.
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                                                                                                                                                                                                                                                                                                                                                   19; Conservative
                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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RESULT 998 AAQ75745/c

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                                                                 Analysis of CDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                            A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0.7%; Score 18.4; DB 1; Length 21;
Best Local Similarity 95.0%; Pred. No. 8.8e+02;
Matches 19; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
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(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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                                                                                         by digestion with restriction enzymes.
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                                                                                                                               Disclosure; Page 9; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                               2708 TAAAAAAAAAAAAAAAA 2727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-018287/03.
                                  WPI; 1995-018287/03
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DB 1; Length 21;

0.7%; Score 18.4;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of ababiled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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               Сарв
                                                                                                                                                                                                                                                                     Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; tive 0; Mismatches 1; Indels
               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
Pred. No. 8.8e+02;
1; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2707 CTAAAAAAAAAAAAAAAA 2726
                                             2709 AAAAAAAAAAAAAAAAA 2728
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Matches 19; Conservative 0;
                                                                         20 AACAAAAAAAAAAAAAA 1
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                                                                                                                                                    AAQ75720 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                             04-AUG-1995 (first entry)
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Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                    Synthetic.
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XX ADK0
XX ADK0
XX O6-M
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KW S8;
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at last once in each nucleic acid and a 3 · variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture computes a sequence of 10-50, particularly 15-55, T residues. The capture of agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-55, T residues. The capture of 10-50, particularly 15-55, T residues. The capture of 10-50, particularly 15-55, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resih, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is spaticularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, or supplements, especially minerals, capture layers of differences in pools of their derivatives, rander of activity of the complete pattern of all mucleic acids from a cell, and can detect very small differences in the nucleic acid solid in a cell, and genes, matrix miniaturisation is based on compatison of nucleic acid pools. Since the capture probes used in the 
                                                                                                                                                                                                                                                                                                                                                                                                       Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                  Boekenkamp D, Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                   (DEGS ) DEGUSSA BIOACTIVES GMBH,
                                                                                                                                                  28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                   28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    and constant regions.
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                                               DE10208794-A1.
                                                                                                04-SEP-2003
  Rattus sp.
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Gaps
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                                     0.7%; Score 18.4; DB-1; Length 21; 95.0%; Pred. No. 8.8e+02; Live 0; Mismatches 1; Indels
Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;
                                                          Local Similarity 95.0
nes 19; Conservative
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2707 CTAAAAAAAAAAAAAAA 2726 20 ò ద

ADK01313 standard; DNA; 21 BP. ADK01313 RESULT 1003 8X4X5X8

Rat DNA microarray capture oligonucleotide #33.

06-MAY-2004 (first entry)

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. 

Rattus sp.

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agants that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sortes of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, computing properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is stranded nucleic acids to the surface is (quasi)covalent, supramolecular strended nucleic acids to the surface is (quasi)covalent, supramolecular is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular is phe method is used (1) for analysis of patterns, especially in mucosal, acids (amino, carboxylic or fatty acid) or through a molecular sieve. The method provides rapid, inexpensive and reproducible captes acids (amino, carboxylic or fatty acid) or their derivatives, or gam call, and can detect very small differences in pools of nucleic acid becomparious of can detect very small differences in the nucleic acid pools, not individual genes, matrix miniaturisation is possible. Ankol281-ADK01344 represent expensed in the method of the invention.

Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

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Gapa ;; 0 0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; 1; Indels 0; Mismatches 19; Conservative Best Local Similarity Query Match Matches

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ADK01319 standard; DNA; 21 BP RESULT 1004 ADK01319

ADK01319; SXB ngs.res

Tue Nov

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable and non-variable regions. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface interaction with streptavidin. The matrix is of plastic, ceramic, glass, interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, thair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxyllic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible can detect very small differences in pools of nucleic acid in a cell, and can lower in page of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid coid or acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                               capture oligonucleotide; pattern; muc cell; food additive; food supplement
                                                                  Rat DNA microarray capture oligonucleotide #39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                              (DEGS ) DEGUSSA BIOACTIVES GMBH.
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                   06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               patterns and screening and constant regions.
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                                                                                                             ss; hybridisation;
blood; nerve; germ
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                                                                                                                                                                                                                                                                                     04-SEP-2003
                                                                                                                                                                                         Rattus sp.
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Sequence 21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; Indels 0; Mismatches Conservative Query Match Best Local Similarity 19; Matches

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4DK01297/c ID ADK01297 standard; DNA; 21 BP.

ADK01297;

(first entry) 06-MAY-2004

mucosal; hair root;

Rat DNA microarray capture oligonucleotide #17.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids acids by bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow binding of particulars sorts of single stranded nucleic acids. The capture organizes are particularly locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceranded nucleic acids to the surface is (LMA) and baving semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b)0)molecular filaments and binding of sangle cranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, surfaced by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, caids (amino, carboxylic or fatty acid) or their derivatives, salts and activity of pharmaceuticals and/or nutritional compounds, e.g. food acids in an ender and on nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible callows inaging of the complete pattern of all nucleic acid sold in the nucleic acid pool; since the method is based on comparison of nucleic acid pool; since the carrive very small differences in the nucleic acid pool; since the carrive very small differences in the nucleic acid pool; the inventor of the inventor of the inventor of the capture probes used in the method of the invention. 

Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;

Gaps ö Query Match
0.7%; Score 18.4; DB 1; Length 21;
Best Local Similarity 95.0%; Pred. No. 8.8e+02;
Matches 19; Conservative 0; Mismatches 1; Indels

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2708 TAAAAAAAAAAAAAAA 2727 | ||||||||||||||||

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Rat DNA microarray capture oligonucleotide #22.
     ADK01302 standard; DNA; 21 BP
                06-MAY-2004 (first entry)
                          ss; hybridisation;
blood; nerve; germ
                                        DE10208794-A1.
                                  Rattus sp.
          ADK01302;
RESULT 1006
   ADK01302
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capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. (DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. 04-SEP-2003.

Dieck HT, Hoppe H; Boekenkamp D,

WPI; 2003-714082/68.

single-stranded nucleic acid, useful for analyzing expression and screening active agents, uses capture agent with variable patterns and screenin and constant regions.

German. Example; Page 5; 8pp; Example; Page 5; 8pp; German.

Comprise Variable anonov region, the capture oligomuciocides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region. That comprises all possible combinations of up to 10 nucleotides to allow binding of particulary sorts of single stranded nucleic acids. The capture of particulary locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (blo)molecular filaments and binding of single conducting stranded nucleic acids to the surface is (quas) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible captor allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual capture probes used in the method of the invention. This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides

Sequence 21 BP; 0 A; 0 C; 3 G; 18 T; 0 U; 0 Other;

Score 18.4; DB 1; Length 21; Pred. No. 8.8e+02; 95.0%; Best Local Similarity Query Match

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                                                                                                                           ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                       Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
  Gaps
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1; Indels
                                                                                                              Rat DNA microarray capture oligonucleotide #37.
0; Mismatches
              2707 CTAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                           Hoppe H;
                       20 CCAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                            (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                  BP.
                                                                                                                                                                                              28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                             28-FEB-2002; 2002DE-01008794.
                                                                  ADK01317 standard; DNA; 21
                                                                                               (first entry)
19; Conservative
                                                                                                                                                                                                                                           3oekenkamp D, Dieck HT,
                                                                                                                                                                                                                                                         WPI; 2003-714082/68.
                                                                                                                                                                 DE10208794-A1.
                                                                                               06-MAY-2004
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                                                                                                                                                   Rattus sp.
                                                                                ADK01317;
Matches
                                                           ADK01317,
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture or comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are biotinylated and immobilised on a surface by capture oligonucleotides are biotinylated and immobilised on a surface capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, c.g. food activity of pharmaceuticals and/or nutritional compounds, or supplements, especially minerals capture probey sentation of differences in pools of mucleic acids from the complete pattern of all mucleic acid benefit and the method is based on complete pattern of all nucleic acid sol, in a cell, and center very small differences

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides of that comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of lo-50, particularly 15-25, T residues. The agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, craticularly a layer of (bio)molecular filaments and expectably in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, carded nucleic acids to the surface is (quasi)covalent, supramolecular, the method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, es, of food activity of pharmaceuticals and/or nutritional compounds, es, of food activity of pharmaceuticals and/or nutritional compounds, es, of food activity of paramaceuticals and/or nutritional compounds, es, of food activity of paramaceuticals and/or nutritional compounds, es, of food activity of paramaceuticals and/or nutritional compounds, es, of food activity of paramaceuticals and/or nutritional compounds, es, of food activity of paramaceuticals and/or nutritional compounds, es, of food mixtures. The method provides rapid, inexpensive and reproducible can detect very small differences in pools of nucleic acids in a cell, allows inaging of t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
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                                                                                                                                                                                                        Gaps
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                                                                                  Query Match
0.7%; Score 18.4; DB 1; Length 21;
Best Local Similarity 95.0%; Pred. No. 8.8e+02;
Matches 19; Conservative 0; Mismatches 1; Indels
Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rat DNA microarray capture oligonucleotide #23.
                                                                                                                                                                                                                                                                                                         2707 CTAAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK01303 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004 (first entry)
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at case in once in each nucleic acid and a 1'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particulars sorts of single stranded nucleic acids. The capture of agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quared nucleic acids to the surface is quasa) covalent, sprond network or garm cells and (ii) for analysis of patterns, especially in mucosal, additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and acids (amino, carboxylic or fatty acid) or their derivatives, salts and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                            Gaps
                                                                                                                                                                                              ö
                                                                                                                                             Length 21;
                                                                                                                                                                                            1; Indels
                                                                                               Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                           Query Match 0.7%; Score 18.4; DB 1; Best Local Similarity 95.0%; Pred. No. 8.8e+02; Matches 19; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat DNA microarray capture oligonucleotide #47.
                                                                                                                                                                                                                                             2707 CTAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                         20 CCAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                            BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                            ADK01327 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dieck HT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DE10208794-A1.
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                                                                                                                                                                                                                                                                                                                                                                RESULT 1009
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mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, a can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
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Sequence 21 BP; 0 A; 2 C; 0 G; 19 T; 0 U; 0 Other;

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Gaps
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 Score 18.4; DB 1; Length 21;
Pred. No. 8.8e+02;
0; Mismatches 1; Indels
                                                   2709 AAAAAAAAAAAAAAAA 2728
                                                                         20 AGAAAAAAAAAAAAAA 1
0.78;
             Local Similarity 95.0 tes 19; Conservative
  Query Match
             Best Loca
Matches
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capture oligonucleotide; pattern, mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                Rat DNA microarray capture oligonucleotide #36.
                                            ADK01316 standard; DNA; 21 BP
                                                                                                                                   (first entry)
                                                                                                                                                                                                                        BB; hybridisation;
blood; nerve; germ
                                                                                                                                      06-MAY-2004
                                                                                         ADK01316;
RESULT 1010
                       ADK01316/
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28-FEB-2002; 2002DE-01008794 28-FEB-2002; 2002DE-01008794 DE10208794-A1. 04-SEP-2003 Rattus sp. 

Dieck HT, Boekenkamp Ď,

(DEGS ) DEGUSSA BIOACTIVES GMBH.

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) there the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) there are in an non-variable regions. The capture oligonucleotides are particular sorts of single stranded nucleic acids. The capture binding of particularly locked nucleic acids (LNA) and the anothor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anothor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quash) capture, supramolecular, physical, stimulated by an electrical field or through a molecular, The method is used (i) for analysis of patterns, especially in mucosal,

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hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cells, and detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
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Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

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Gaps
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0.7%; Score 18.4; DB 1; Length 21;
95.0%; Pred. No. 8.8e+02;
ve 0; Mismatches 1; Indels
                                                    2708 TAAAAAAAAAAAAAAAA 2727
                                                                     20 TGAAAAAAAAAAAAAA 1
             95.0%;
                           Conservative
            Local Similarity
nes 19; Conserv
  Query Match
               Best Loca
Matches
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ВР ADK01299 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01299; RESULT 1011 ADK01299/c

capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. Rat DNA microarray capture oligonucleotide #19. blood; nerve; germ ss; hybridisation;

28-FEB-2002; 2002DE-01008794. DE10208794-A1. 04-SEP-2003. Rattus sp. 

(DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794.

Hoppe Boekenkamp D, Dieck HT,

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WPI; 2003-714082/68.

useful for analyzing expression uses capture agent with variable Sorting single-stranded nucleic acid, patterns and screening active agents, and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid bools, then reading out, where the nucleic acids are selectively bound using appture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface

is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, by selectrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, chair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent 

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

Gaps ö Query Match

0.7%; Score 18.4; DB 1; Length 21;
Best Local Similarity 95.0%; Pred. No. 8.88+02;
Matches 19; Conservative 0; Mismatches 1; Indels 2708 TAAAAAAAAAAAAAAA 2727 20 TCAAAAAAAAAAAAAAA 1 ઠે g

BP ADK01301 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01301; RESULT 1012 ADK01301/c 

Rat DNA microarray capture oligonucleotide #21.

capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. ss; hybridisation; blood; nerve; germ

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

30ekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agente that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5' invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3' variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow agents are particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The

capture original receptantion. The matrix is of plastic. Ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, chir root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible cailows inaging of the complete pattern of all nucleic acid in a cell, and can lows inaging of the complete pattern of all nucleic acid in a cell, and can matrix minaturisation of nucleic acid pool. Since the method is based on comparison of nucleic acid pools. Since the method is based on comparison of since nice acid pools. Allows in a cell, mathod in a possible. AbRO1281-AbRO1344 represent capture oligonucleotides are biotinylated and immobilised on a surface by capture probes used in the method of the invention. 

Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other;

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Gaps ö Score 18.4; DB 1; Length 21; Pred. No. 8.8e+02; 1; Indels 0; Mismatches 0.7%; 95.0%; Query Match
Best Local Similarity 95.0
Matches 19, Conservative

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ADK01315 standard; DNA; 21 06-MAY-2004 (first entry) ADK01315; RESULT 1013 

ВЪ.

Rat DNA microarray capture oligonucleotide #35.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5' invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region

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that comprises all possible combinations of up to 10 nuclectides to allow binding of particular sorts of single stranded nucleic acids. The capture of agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceracity resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single cranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, cramethod is used (i) for analysis of patterns, especially in mucosal, acity to pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acidis (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible con additives of additives of additives of allows inaging of the complete pattern of all nucleic acid from cells. It callows inaging of the complete pattern of all nucleic acid bool. Since the conduction is possible. ADROIS41-ADROI344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 capture probes used in the method of the invention.
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0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; Live 0; Mismatches 1; Indels Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other; Best Local Similarity Query Match

19; Conservative

Matches

2708 TAAAAAAAAAAAAAAA 2727 20 TGAAAAAAAAAAAAA 1 à 셤

ADK01326 standard; DNA; 21 BP ADK01326; ADK01326/ 

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #46.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture

degenes interdate (af immodified on the surface of a surface of agenes interdate and non-variable end as 3 -variable, discriminatory region. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particulary locked nucleic acids (LMA) and the anchor region comprises are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, certain gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, computed by an electrical field or through a molecular, cativity of pharmaccuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acidis (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible callows inaging of the complete pattern of all nucleic acids from cells, and can lows inaginal differences in the nucleic acid pool. Since the callows inaginal matrix anniaturisation is possible, AMROGI281-AMROGI344 represent conversent elements. agents that are (a) immobilised on the surface of a solid matrix and (b) capture probes used in the method of the invention. 8×999999999999999999999999

Sequence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;

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Gaps ; 0

Gaps ö 0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; artive 0; Mismatches 1; Indels Conservative Similarity 19; Query Match Best Local 8 Best Loca Matches

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8 셤 RESULT 1015

BP ADK01300/c ID ADK01300 standard; DNA; 21

ADK01300;

(first entry) 06-MAY-2004

Rat DNA microarray capture oligonucleotide #20.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

Hoppe H; Boekenkamp D, Dieck HT,

(DEGS ) DEGUSSA BIOACTIVES GMBH.

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions. 

Example; Page 5; 8pp; German.

Tue Nov

20 TCAAAAAAAAAAAAAAAA 1 RESULT 1016 Matches g ઠે

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The capture oligonucleotides have a 5-invariable anchor region. The complement of which is present at least once in each mucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises are particularly locked nucleic acids (LNA) and the anchor region comprises as sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b)omolecular filaments and binding of shingle cranded nucleic acids to the surface is quasi) covalent, supramolecular, physical, surmulated by an electrical field or through a molecular sieve. The method is used (1) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional constructions or supplements, especially minerals, trace elements or store and activity of pharmaceuticals in pools of nucleic acids from cell, and each or comparison of nutritional cacid section of additives and an each or supplements and the nucleic acid sold in the marthy of the individual can capture probes used in the method of the invention

/ Match 0.7%; Score 18.4; DB 1; Length 21; Local Similarity 95.0%; Pred. No. 8.8e+02; les 19; Conservative 0; Mismatches 1; Indels Seguence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other; 2708 TAAAAAAAAAAAAAAA 2727 Query Match

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Gaps

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ADK01310 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01310;

Rat DNA microarray capture oligonucleotide #30.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

DE10208794-A1 04-SEP-2003 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. (DEGS ) DEGUSSA BIOACTIVES GMBH

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression

Boekenkamp D, Dieck HT, Hoppe H;

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using apprure caging out, where the nucleic acids are selectively bound using apprure agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow binding of particulars sorts of single stranded nucleic acids. The capture of agents are particularly locked nucleic acids (LNA) and the anchor region that comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (b) molecular filaments and binding of anigle cranded nucleic acids to the surface is (quasi) covalent, supramolecular or physical, surmulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, thair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic crais (main, oraboxylic or fatty acid) or their derivatives, salts and acids in an end or acids from cells. It allows inaging of the complete pattern of all nucleic acids from cell; and crandetect very small differences in the nucleic acid soil and in a cell, and crandet and an acid pool. Since the cannow may be matrix and in the method is possible, the interior or detect very small differences in the interior acid pool. Since the converse and an enterior or the presention of the method is possible. The interior or detect very small difference in the interior or detect very small miniature patterns and screening active agents, uses capture agent with variable capture probes used in the method of the invention. Example; Page 5; Bpp; German. and constant regions. 

ö 0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; 1; Indels Sequence 21 BP; 0 A; 0 C; 2 G; 19 T; 0 U; 0 Other; 95.0%; Pred. .v. 19; Conservative Local Similarity Query Match Matches

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Gaps

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Rat DNA microarray capture oligonucleotide #31. ADK01311 standard; DNA; 21 BP. (first entry) 06-MAY-2004 ADK01311; ADK01311/

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement. (DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. DE10208794-A1. 04-SEP-2003. Rattus sp. 

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture care dependent are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (IMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, certain, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface of spraticularly a layer of (b)0)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, acitives or supplements, especially minerals, trace elements, organic acids and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (mino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible mixtures. The method provides rapid, inexpensive acid in a cell, and can lows inaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual computed in the method is possible, havened acid pools. Since the capture and an acuter provibes used in the method is passible, hav
                                                                                     Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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                                                                                                                                                                                                  Example; Page 5; 8pp; German.
                                                                                                                                             and constant regions
                             WPI; 2003-714082/68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-APR-2002
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Matches
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Gaps . 0 0.7%; Score 18.4; DB 1; Length 21; 95.0%; Pred. No. 8.8e+02; 1; Indels 0; Mismatches 2706 ACTAAAAAAAAAAAAAA 2725 20 ACAAAAAAAAAAAAAA 1 95.08; 19; Conservative Local Similarity

PolyA adaptor oligonucleotide SEQ ID NO:1. BP ABA93238 standard; DNA; 22 (first entry) 

Detection; comparative detection; adaptor; ss.

Synthetic.

JP2001333800-A.

04-DEC-2001.

30-MAY-2000; 2000JP-00160324.

30-MAY-2000; 2000JP-00160324

The present invention describes a method for the comparative detection of the amount of an RNA. The method comprises: (a) cDNAs obtained by transcribing respectively from at least two tissue RNAs are respectively from at least two tissue RNAs are respectively from at least two tissue RNAs are respectively fragmented by using a same restriction enzyme; (b) each different adaptor and a common adaptor are added to each of the cDNA fragments derived from adad cDNAs are mixed together; (d) an adaptor primer having the common sequence to said different adaptor and a gene specific adaptor are used to amplify said adaptor-added cDNAs containing no region derived from polyadenylic acid of the mRNA before the addition of the adaptor among the measured between the tissues; (f) the RNA is detected from the measured result; (g) each different adaptor and a common adaptor of the adaptor acid of the genomic DNA fragments derived from a same or different individuals; (h) the resultant adaptor-added genomic DNAs are mixed together; (i) the daptor-added genomic DNAs are applified by using an adaptor primer having the common sequence to the different adaptor and adaptor and adaptor and deprecent of the amounts of the amplified amounts of the genomic DNAs are measured between the individuals. The method is cused for the detection of the amounts of RNA and DNA. The present core exemplification of the present invention Comparative detection of the amounts of RNA and DNA. Disclosure; Page 9; 9pp; Japanese. (UNIT-) UNITECH CO LID. WPI; 2002-135950/18. 

0.7%; Score 18.4; DB 1; Length 22; 95.0%; Pred. No. 9e+02; ive 0; Mismatches 1; Indels Sequence 22 BP; 19 A; 1 C; 1 G; 1 T; 0 U; 0 Other; Query Match 0.73 Best Local Similarity 95.0 Matches 19; Conservative

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Gaps

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3 TCAAAAAAAAAAAAAA 22 ઠે 셤

AAQ75028 standard; DNA; 23 BP.

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RESULT 1019

AAQ75028;

Synthetic oligo; solid phase immunoassay; ss. 94WO-US005407 (first entry) (revised) WO9426932-A1 13-MAY-1994; 24-NOV-1994. 25-MAR-2003 03-AUG-1995 LCR oligo 2. Synthetic

(USSH ) US DEPT HEALTH & HUMAN SERVICES. 93US-00061694. Khudyakov YE; 13-MAY-1993; Fields HA, 

WPI; 1995-006819/01.

Solid phase immunoassay using oligo:nucleotide as label - also new conjugates of oligo:nucleotide coupled to antigenic peptide, partic. for

diagnosing hepatitis C or E virus infection.

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AAR62941 and AAR62942 are examples of synthetic immunoreactive peptides. They are used in a method for detecting an antigen in a subject. The method involves binding the antigen to a solid support and then reacting it with an immunoreactive ligand (L) bound to an oligo; removing any unreacted L, and then detecting the presence of the oligo. A similar method can be used to detect Abs, in which case the ligand is an oligo-labelled Ag. The use of an amplifiable oligo as the label allows Ag or Ab to be detected at very low levels. An exemplary olgi is AAQ75024 which can be covalently attached by the 5' terminus to the N- or C-terminal of a synthetic peptide. For LCR using oligo AAZ75024, oligos 1-4 (see
                                                             AAR62941 and AAR62942 are examples of synthetic immunoreactive peptides. They are used in a method for detecting an antigen in a subject. The method involves binding the antigen to a solid support and then reacting it with an immunoreactive ligand (L) bound to an oligo; removing any unreacted L, and then detecting the presence of the oligo. A similar method can be used to detect Abs, in which case the ligand is an oligolabelled Ag. The use of an amplifiable oligo as the label allows Ag or Ab to be detected at very low levels. An exemplary olg is AAQ75024 which as synthetic peptide. For LCR using oligo AAZ75024, oligos 1-4 (see AAQ75027-Q75030) can be used. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Solid phase immunoassay using oligo:nucleotide as label - also new conjugates of oligo:nucleotide coupled to antigenic peptide, partic. for diagnosing hepatitis C or E virus infection.
                                                                                                                                                                                                                                                                                                                      0.7%; Score 18.4; DB 1; Length 23; Local Similarity 95.0%; Pred. No. 9.2e+02; Conservative 0; Mismarrh.
                                                                                                                                                                                                                                                                                                              Seguence 23 BP; 19 A; 4 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic oligo; solid phase immunoassay; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 13; 34pp; English.
                                  Example, Page 13; 34pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ75029 standard; RNA; 23
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(first entry)
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03-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated limonene hydroxylase nucleic acids - which encode limonene-6-hydroxylase and limonene-3-hydroxylase, which can be used to produce trans-carveol and trans-isopiperitenol.
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                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                    (-)-limonene-6-hydroxylase; (-)-limonene-3-hydroxylase; L3H; L6H; spear mint; peppermint; enzyme; limonene hydroxylase; trans-carveol; trans-isopiperitenol; pathogen defense mechanism; attractant; environmental signal; monoterpene hydroxylase; PCR primer; ss.
 correct PN
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                                                                                   Length 23;
AAQ75027-Q75030) can be used. (Updated on 25-MAR-2003 to
                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                BP; 0 A; 0 C; 4 G; 1 T; 18 U; 0 Other,
                                                                                 Score 18.4; DB 1;
Pred. No. 9.2e+02;
                                                                                                                     0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                       (-)-limonene-6-hydroxylase primer 3.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; Page 27; 80pp; English.
                                                                                                                                                     2707 CTAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                      20 CCAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                         AAX06572 standard; DNA; 19 BP.
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                                                                                                    95.0%;
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Matches 18; Conservative
                                                                                                                     19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lupien SL,
                                                                                                      Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Mentha spicata.
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                                                     Sequence 23
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Fatty lesion development, atherosclerosis, Alzheimer's disease, nervous system disorder; Parkinson's disease; immune system disorder; stokaemia, lymphopaenia, leukocyte adhesion deficiency syndrome; haemoglobinuria, anaemia, hyperproliferative disorder; gucher's disease; coagulation disorder; blood platelet disorder; autoimmune disorder; dermititis, herpes simplex; Addison's disease; rheumatoid arthritis;

3' sequencing primer #1 to identify and characterise polynucleotides

(first entry)

01-NOV-2001

AAD15201;

Grave's disease; gene therapy; antiarteriosclerotic; immunostimulant; cardiovascular; antiviral; primer; ss.

AAD15201 standard; DNA; 19 BP.

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The present primer was used to reverse transcribe cDNA encoding a C-20 oxidase. The amplifie fragment is used in the method of the invention. The specification describes methods for the inhibition and control of gibberellic acid levels. Gibberellic acid levels may be inhibited or controlled by use of a chimeric expression construct expressing a RNA protein which suppresses the gibberellin biosynthetic pathway sequence, diverts substrate from the pathway, or degrades pathway substrates or products. The methods uses copalyl diphosphate synthase, is beta-hydroxylase, 2-oxidase, phytoene synthase, C-20 oxidase, and a lateral phytosne synthase, c-20 oxidase, and a lateral phytosne synthase, c-20 oxidase, and a lateral phytosne synthase, c-20 oxidase, and can seed germination and seedling growth especially to regulate gene products of gibberellin biosynthetic pathway and restoration of normal seed germination, in transgenic plants. The plants products are gibberellin deficient, and have shortened hypocotyl and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Obtaining transgenic plant useful for controlling seed germination and seedling growth comprises transgene comprising a sequence expressing altered levels of an essential hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kishore GM, Logusch EW, Logusch SJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gibberellic acid, copalyl diphosphate synthase, 3beta-hydroxylase, 2-oxidase; phytoene synthase, C-20 oxidase, 2beta,3beta-hydroxylase, seed germination; seedling growth; gibberellin biosynthetic pathway; transgenic plant; hypocotyl; epicotyl; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer HOOK for cDNA encoding a C-20 oxidase polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epicotyl phenotypes compared to normal plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 17; Page 262; 267pp; English.
Heck GR,
                                                                                                                                                                                                                                                            ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-US018066.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98US-0096111P.
99US-0137977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rao S, Ream JE;
                                                                                                                                                                                                                                                AAZ99489 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                               03-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Elich TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MONS ) MONSANTO CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-224351/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cucurbita maxima.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200009722-A2.
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07-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brown SM, F
Piller KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-FEB-2000
                                                                                                                                                                                                                                                                                                                            AAZ99489;
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   2708
                                                                                                                                                                                                                      AAZ99489/
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New polynucleotides regulated by fatty lesion development and their encoded polypeptides, useful for preventing, treating or ameliorating atherosclerosis, as well as for immune or hyperproliferative disorders.

Example 1; Page 79; 188pp; English.

Hasel KW;

Glass JR, Sutcliffe JG,

(DIGI-) DIGITAL GENE TECHNOLOGIES INC.

Sartani A,

Leonardi A,

WPI; 2001-514526/56.

25-JAN-2001; 2001WO-US002439. 25-JAN-2000; 2000US-0177963P.

WO200154651-A2. Unidentified.

02-AUG-2001.

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fatty lesion development, which comprises any of 55 polynucleotide sequences from Oryctolagus cuniculus. The polynucleotide, polypeptide or antibody is useful for preventing, treating, modulating or ameliorating a medical condition, particularly atherosclerosis. The invention is used as marker or detector of nervous system disorder or disease (e.g. parkinson's disease, Alzheimer's disease, ischaemia, dementia). The invention may also be useful for treating deficiencies or disorders of the immune system (e.g. lymphopaenia, leukocyte adhesion deficiency syndrome or haemoglobinuria, anaemia), hyperproliferative disorders (e.g. Gaucher's disease), infectious disease (e.g. herpes simplex), coagulation disorders, blood platelet disorders and autoimmune disorders (Addison's disease, rheumatoid arthritis, dermatiis, Grave's disease). The polynucleotide sequence is a lso used in gene therapy. The present sequence is a 3' sequencing primer used in the identification and characterisation of polynucleotides up-regulated by fatty lesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to an isolated nucleic acid regulated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 18.2; DB 1;
Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2708 TAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH21968 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   development
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19
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ID AAH2
XX
AC AAH2
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Gaps . 0

0; Indels

DB 1; Length 19;

Score 18.2; DB 1; Pred. No. 8.6e+02; 1; Mismatches 0

0.7%;

2708 TAAAAAAAAAAAAAA 2726

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Conservative

Local Similarity

Query Match Best Loca Matches

18;

BAAAAAAAAAAAAA 1

13

RESULT 1023 AAD15201/c

ngs.res

(first entry)

AAF76617;

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digital sequence tag; expressed sequence tag; neuroleptic; antimanic; central nervous system; antidepressant; gene therapy, diagnosis; neuropsychiatric disorder; schizophrenia; bipolar disorder; addiction-related behaviour; chromosome identification; immune response;
                           Mouse total gene expression analysis (TOGA) 3' sequencing primer SEQ:92.
                                                                                                                                                                                                                                                                                  New neuroleptic-regulated polynuclectides expressed in the central nervous system for diagnosing and treating neuropsychiatric disorders such as schizophrenia, bipolar disorder and addiction-related behavior.
                                                human; total gene expression analysis; TOGA; DST; EST
                                                                                                                                                                                                                           (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                               Example 1; Page 87; 210pp; English
                                                                                                                                                                                 26-OCT-2000; 2000WO-US029690.
                                                                                                                                                                                                      99US-0161379P.
                                                                                                                                                                                                                                              Sutcliffe JG,
         (first entry)
                                                                                                   PCR primer; probe; ss.
                                                                                                                                                                                                                                                                  WPI; 2001-300499/31.
                                                                                                                                          WO200130972-A2.
                                                                                                                                                                                                       26-OCT-1999;
                                                                                                                       musculus
         16-AUG-2001
                                                                                                                                                                03-MAY-2001.
                                                                                                                                                                                                                                               Thomas EA,
                                                                                                                        MuB
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The present invention describes isolated neuroleptic-regulated nucleic acid molecules. (I) have neuroleptic, antimanic and antidepressant acid molecules. (I) have neuroleptic, antimanic and antidepressant activities, and can be used in gene therapy. (I), polypeptides (II) encoded by (I), or a host cell (III) comprising (I), are useful for preventing, treating, modulating or ameliorating a medical condition such as a neuropsychiatric disorder. (I) are useful as diagnostic agents for diagnostic a pathological condition such as neuropsychiatric disorder e.g. schizophrenia, a bipolar condition such as neuropsychiatric disorder e.g. schizophrenia, a bipolar disorder or addiction-related behaviour. (I) are useful for detecting the control gene expression through triple helix formation ample. (I) can be used as probes and primers, for chromosome control gene expression through triple helix formation identification, to control gene expression through triple helix formation control gene expression through triple helix formation disorders, identifying individuals from minute biological samples, as an alternative to restriction fragment length polymorphism (RFLP) and as polymorphic markers for forensic purposes. (I) is also useful as molecular weight markers on Southern gels, diagnostic probes for the presence of specific mRNA in a particular cell type, as a probe to subtract-out known sequences in the process of discovering novel controlledides, for selecting and making oligomers for attachment to a gene chip or other support, to raise anti-DNA antibodies using DNA immunisation technique, and as an antigen to elicit an immune response. The areas an antigen to elicit an immune response. The areas an antigen to elicit an immune sequences used in the areas and antigen to elicit an immune sequence. Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other; invention exemplification of the present

2; DB 1; Length 19; 8.6e+02; 0.7%; Score 18.2; Di 94.7%; Pred. No. 8.6e iive 1; Mismatches Query Match Best Local Similarity Matches 18; Conserv

2708 TAAAAAAAAAAAAAA 2726 BAAAAAAAAAAAAAAA 1 13 g ઠ

18; Conservative

AAF76617 standard; DNA; 19 BP

ö The present invention provides the protein and coding sequences of the peppermint and spearmint (-)-limonene-3-hydroxylase and the spearmint (-)-limonene-f-hydroxylase and pCR primers which were used to isolate the sequences. These are useful in the production of transgenic plants with altered flavour and aroma Novel isolated limonene hydroxylase encoding nucleic acid molecule, useful for altering production of limonene-6-hydroxylase or limonene-3-Gaps ô Spearmint; peppermint; (-)-limonene-6-hydroxylase; (-)-limonene-3-hydroxylase; flavour; aroma; probe; PCR primer; Score 18.2; DB 1; Length 19; pred. No. 8.6e+02; Spearmint (-)-limonene-6-hydroxylase PCR primer SEQ ID NO: 0; Indels Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other; 1; Mismatches (UNIW ) UNIV WASHINGTON STATE RES FOUND. hydroxylase in suitable host cell. Example 4; Col 55; 57pp; English ۲. ن Karp 0.7%; 97US-00881784. 99US-00292768 18; Conservative Croteau RB, Lupien SL, Query Match Best Local Similarity WPI; 2001-243405/25. Mentha spicata US6194185-B1. 14-APR-1999; 24-JUN-1997; 15-MAY-2001 27-FEB-2001. Matches 8

Hasel KW;

Pribyl TM, Hilbush B,

Mouse microglia and macrophage regulatory gene primer #60. 2708 TAAAAAAAAAAAAAAA 2726 19 DAAAAAAAAAAAAAA 1 BP. AAS06525 standard; DNA; 19 (first entry) 07-SEP-2001 AAS06525; RESULT 1026 셤

Mouse, microglia, macrophage; regulatory gene; digital sequence tag; DST; PCR-based total gene expression analysis; TOGA, infectious disorder; neuroinflammatory pathology; neurodegenerative disease; gene therapy; hyperproliferative disorder; autoimmune; inflammatory disorder; primer; Mus musculus.

06-NOV-2000; 2000WO-US030585. 17-MAY-2001

WO200134770-A2

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Gaps

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Indels

Tue Nov

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The present sequence represents a primer used to isolate novel mouse microglia and macrophage regulatory gene DST (digital sequence tag) sequences. A806401-A806590 represent these novel sequences and the sequences used to isolate them. The PCR-based total gene expression analysis (TOGA) system is used to examine the expression pattern of molecules corresponding to genes that are regulated in cartivated microglia, activated microglia, unstimulated macrophage and activated macrophage. The polymolecities of the invention, the polympetides encoded by them and antibodies that bind to these polympetides are useful are useful for the diagnosis, prevention, treatment or amelioration of a medical condition, preferably a neuroinflammatory pathology or a neurodegenerative disease such as Alzeimer's disease, senile dementia, Parkinson's disease, obsessive compulsive disorders, epilepsy, schizophrenia, multiple sclerosis, depression and bipolar manic-depressive disorders (e.g. severe combined immunodeficiency, SCID) cancer), immune disorders (e.g. severe combined immunodeficiency, SCID) cancer), immune disorders (e.g. insulin dependent diabetes mellitus).
                                                                                                                                                                      New regulated genes of microglia and macrophages, useful for diagnosing, preventing or treating neuroinflammatory pathology and neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Central nervous system; CNS; neuroleptic; mouse; human; psychoses; neuropsychiatric disorder; psychiatric disorder; Alzheimer's disease; Pick's disease; Binswanger's disease; sencephalopathy; Parkinson's disease; obsessive compulsive disorder; epilepsy; ischaemia; addiction; multiple sclerosis; depression; manic-depressive disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 18.2; DB 1; Length 19;
Pred. No. 8.6e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                               Tobal GM;
                                                                                                           Sutcliffe JG, Almazan MT,
                                                                      (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                             Example 1; Page 88; 244pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2708 TAAAAAAAAAAAAAA 2726
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99WO-US026824.
                 03-MAR-2000; 2000US-0186770P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 94.7'
Matches 18; Conservative
                                                                                                                                               WPI; 2001-308782/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy
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12-NOV-1999;
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                                                                                                             Carson MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; ss
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This invention relates to the CDNA sequences of novel isolated polynucleotides associated with psychoses or other neuropsychiatric consistences of the invention may act as blockers of D. 2 receptors in the meso-limbic dopamine system. The nucleotide sequences of the invention and the polypeptides encoded by them are useful in the meso-limbic anedical condition e.g. a neuropsychiatric disorder. An antibody that binds the proteins of the invention is useful for antibody that binds the proteins of the invention is useful for preventing, treating, modulating or ameliorating neurological disorders or antibody that binds the proteins of the invention is useful for such as psychoses or other neuropsychiatric disorders or a susceptibility to a neurological disorder such as psychoses or other neuropsychiatric disorders or a susceptibility to a neurological disorder such as psychoses or other neurological disorders and other containing the alteration (increase or decrease) in the expression of determining the alteration (increase or decrease) in the expression of deficiencies or disorders of the invention are useful in treating deficiencies or disorders of the central nervous system or peripheral correction or mobilisation (chemotaxis) of neuroblasts, stem cells or glial cells The sequences are useful as a marker or detector of a corricular nervous system fisease or disorder such as Alzheimer's disease, pick's disease, Binswanger's disease, other senile dementia, particular nervous system compulsive disorders, epilepsy, encephalopathy, ischaemia, addiction, multiple sclerosis, depression and manic-depressive disorder. The present sequences are setulation of the containing the province of a collgonucleotide primer used in the identification of the containing the colls of the containing the province or disconders and collgonucleotide primer used in the invention of the containing the collsonal and the containing the province or disconders.
                                                                                                                                                                                                                   New polynucleotide useful in gene therapy for preventing, treating modulating or ameliorating a medical condition such as psychoses or a neuro psychiatric disorder e.g. schizophrenia, or a bipolar disorder in a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rabbit; Oryctolagus cuniculus; atherosclerosis; intimal hyperplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                      Thomas EA, Sutcliffe JG, Pribyl TM, Hilbush BS, Hasel KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 18.2; DB 1; Length 19; 94.7%; Pred. No. 8.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
                                                                                        (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 40; 254pp; English.
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                     29-SEP-2000; 2000US-0236790P.
18-JAN-2001; 2001US-0263084P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 94.7
nes 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryctolagus cuniculus.
                                                                                                                                                                                    WPI; 2002-383271/41.
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Gaps ; 0

01-OCT-2001; 2001WO-US030695.

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Hepatitis B virus; HBV infection; chronic hepatitis; toxicity; virucide; acute hepatitis; therapeutic; gene therapy; vaccine; infectious disease; TOGA; TOtal Gene expression Analysis; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                      19 BAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                        AAD34663 standard; DNA; 19 BP
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                  21-NOV-2001; 2001WO-US044072
                               21-NOV-2000; 2000US-0252216P
                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                              Conservative
                                                         Sartani A,
                                                                                                                                                                                                                                                       the present invention
                                                                                                                                                                                                                                                                                     Local Similarity
es 18; Conserv
                                                                      WPI; 2002-575233/61
                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200222783-A2
                                                                                                                                                                                                                                                                                                                                                                                   16-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAR-2002
                                                           Leonardi A,
                                                                                                                                                                                                                                                                                                                                                                      AAD34663
                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                            RESULT 1029
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The present invention describes an isolated polymucleotide (I) and its complements, and degenerate variants, comprising a sequence selected from complements, and degenerate variants, comprising a sequence selected from those given in ABO73226 to ABO73222 (NS), which is a digital sequence tag (DST) corresponding to mRNAs whose expression is regulated by proliferative lesion development caused by mechanically induced intimal proliferative lesion development creatment, or by proliferative lesions and reversed by lercanidipine treatment, or by proliferative lesions can deversed by lercanidipine treatment. (I) has antiatherosclerotic activity and can be used in gene therapy. (I) can be used for diagnosing cativity and can be used in gene therapy. (I) can be used for diagnosing the medical condition hased on the presence or absence of the mutation. (I) is also useful for diagnosing atherosclerosis, or the susceptibility continuous conditions and also condition such as antherosclerosis in amount of expression of (I). (I) is also used for preventing, treating, modulating, or ameliorating a medical condition such as atherosclerosis in amammalian subject. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polynucleotides related to regulated genes characteristic of atherosclerosis, useful for diagnosing, preventing, treating, moc ameliorating atherosclerosis in a mammalian subject.
                                                                                                                                                                                                                                                                                                                                                                                                             Glass JR, Hasel KW;
                                                                                                                                                                                                                                                                                                  (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 28; 130pp; English
30-MAY-2002
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The present invention relates to a method for preventing, treating, modulating or ameliorating a medical condition. The method involves modulating one or more nucleic acid molecules up- or down-regulated in hepatitis B virus (HBV) infection or polypeptides encoded by the nucleic acid molecules or antibodies that bind to the polypeptide. The method is useful for preventing, treating, modulating or ameliorating a medical condition. It is also useful for determining the presence or absence of a mutation in the nucleic acid molecules or detecting an alteration in expression of the polypeptide which is useful for the diagnosis of the patitis viral infection (e.g., acute hepatitis versus chronic hepatitis viral infection and a gene expression publication and a gene expression profile is useful for identifying polypeptides and polynucleotides which are associated with charaptitis viral infection. Sequences of the invention are used in gene therapy, and as vaccines. Nucleic and sequences are useful as a

Preventing and treating hepatitis viral infection in a mammal, comprises administering nucleic acid molecules that up- or down-regulate in hepatitis B virus infection or polypeptides encoded by the nucleic acid

Disclosure; Page 28; 125pp; English.

molecules.

Hilbush BS

Mueller R,

Guidotti LGDVM,

SF,

Wieland

₹,

Chisari

WPI; 2002-339865/37.

DIGI-) DIGITAL GENE TECHNOLOGIES INC.

15-SEP-2000; 2000US-0233176P.

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Gibberellin, transgenic plant, seed germination, seedling growth, GA,
transgenic, 2beta-3beta hydroxylase, enzyme, pumpkin, PCR, primer; ss
                                                                                                                                                                                             HOOK PCR primer used to isolate pumpkin 2beta-3beta hydroxylase cDNA.
0.7%; Score 18.2; DB 1; Length 19; 94.7%; Pred. No. 8.6e+02;
                      0; Indels
                        1; Mismatches
                                              2708 TAAAAAAAAAAAAAA 2726
                                                               BP.
             94.78;
                                                                                                                              AAD40279 standard; DNA; 19
                                                                                                                                                                           (first entry)
                          18; Conservative
              Local Similarity
                                                                                                                                                                                                                                                                                 US2002053095-A1.
                                                                                                                                                                                                                                                            Cucurbita pepo.
                                                                                                                                                                           22-OCT-2002
                                                                                                                                                    AAD40279;
   Query Match
                                                                                                         1030
                          Matches
                                                                                                        RESULT 10
AAD40279/
                                                                                                                                g
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                                                                                                                                                                                                                                                 PCR primer #4 used for direct sequencing of TOGA generated PCR products.
                                                                          Gaps
                                                                          .;
0
                                                  DB 1; Length 19;
                                                                         0; Indels
                            Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                   0.7%; Score 18.2; DB 1;
94.7%; Pred. No. 8.6e+02;
ative 1; Mismatches 0
                                                                          1;
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Kishore GM, Logusch EW, Logusch SJ;

Heck GR,

Brown SM, Elich TD,

(BROW/) BROWN S M.

10-AUG-1999; 10-AUG-1999;

99US-00371307. 99US-00371307.

02-MAY-2002.

ö

Gaps

ö

generated PCR

diagnostic markers for HBV infection and for treating infectious diseases. The present DNA sequence is a PCR primer which is used for direct sequencing of TOGA (TOtal Gene expression Analysis) generated

Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;

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Gaps

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Example 10; Page 54; 85pp; English.

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The invention relates to control of gibberellin (GA) levels in plants.

The method involves producing transgenic plants having a phenotype of reduced seed germination and reduced early seedling growth, then restoring seed germination and early seedling growth by treating plants with an appropriate compound when conditions are favourable. The method is useful to control seed germination and/or early seedling growth in ggricultural production so that unfavorable environmental conditions normally reducing agronomic output can be avoided and yields increased. Plants also demonstrate increased uniformity of germination, emergence and seedling vigor, so increasing yields at harvest. The method is especially useful in crop plants such as e.g. canola, soybean, cotton, cotto, and is also useful in storage and transport of seeds to reduce premature germination which may affect agronomic or food quality of the seeds. The present sequence is a PCR primer used to isolate pumpkin 2beta in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New chimeric polypeptide comprising a histone acetyltransferase polypeptide segment and a segment comprising a histone deacetylase chromatin-associated protein complex subunit, useful for modulating gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Histone acetyltransferase; histone deacetylase; gene expression profile; chromatin-associated protein; gene expression; primer; ss.
                                                                    Control of gibberellin levels in plants useful to avoid unfavorable conditions in crops to increase yields, using transgenic plants having reduced seed germination and early seedling growth then treatment to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reverse transcription primer used to produce yeast cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 18.2; DB 1;
94.7%; Pred. No. 8.6e+02;
vative 1; Mismatches 0;
                                                                                                                                                           Example 19; Page 104; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2708 TAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ68389 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-JUN-2002; 2002WO-US019750.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-JUN-2001; 2001US-0300135P.
   Ream JE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                           restore these properties.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-175280/17.
                                   WPI; 2002-489107/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression in cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
 Rao S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CERE-) CERES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003000715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-APR-2003
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 Piller
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polypeptides. The chimeric polypeptides comprise a polypeptide segment that exhibits histone acceyltransferase activity, and a polypeptide segment that exhibits histone acceyltransferase activity, and a polypeptide segment having 40% or greater sequence identity to a subunit of a histone deacetylase chromatin-associated protein complex. The chimeric polypeptides are useful for determining gene expression profiles in specific cells, for modulating gene expression in specific cells, and for making genetically modified enkaryotes. The present sequence represents a reverse transcription primer used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pathological condition (comprises administering to a mammalian subject at least one of: (a) a first polynucleotide comprising a sequence having at least one of: (a) a first polynucleotide comprising a sequence having 38-889 bp (consisting of the sequences in ACC79319 to ACC7932 (I)) or a second polynucleotide at least 95* identical to the first polynucleotide; (b) a third polynucleotide comprising at least 10-bp sequence that is hybridisable to the first polynucleotide under stringent conditions; or (c) a gene corresponding to any of (1)-(2) or another gene at least 95* identical to the gene. (I) have cytostatic activities, and can be used in vaccines and in gene therapy. The method is useful for preventing or treating e.g., ataxia telanglectasia (AT), AT tumours or other cancers. ACC79393 to ACC793423 represent primers used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Preventing or treating a pathological condition e.g., ataxia telangiectasia (AT), AT tumors or other cancers comprises administering polynucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method for preventing or treating a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pathological condition; ataxia telangiectasia; AT; tumour; cancer; cytostatic; vaccine; gene therapy; PCR primer; ss.
                                       specification describes chimeric histone acetyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Callahan MLA, Pankratz DG, Vibat CRT;
                                                                                                                                                                                                                                                                                0.7%; Score 18.2; DB 1; Length 19;
94.7%; Pred. No. 8.6e+02;
ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M13 sequencing primer 3' primer SEQ ID NO:84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 76; 184pp; English.
                                                                                                                                                                                                                                                                                                                                                                2708 TAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-OCT-2002; 2002WO-US033311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-OCT-2001; 2001US-0330206P.
                                                                                                                                                                                                                                                                                                     94.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACC79402 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enterobacteria phage M13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Winrow CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-393520/37.
                                                                                                                                                                                                                                                                                                   Local Similarity
nes 18; Conserv
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Warren AJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACC79402;
                                                                                                                                                                                                                                                                                Query Match
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Matches
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Tue Nov

SXS

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BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;

Seguence 19

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Atherosclerosis, vaccine, nervous system disorder, Alzheimer's disease, Parkinson's disease, multiple sclerosis, immune disorder; gene therapy, autoimmune disorder; rheumatoid arthritis, hyperproliferative disorder; haemolytic namemia; graft-versus-host disease; inflammation; infection; epilepsy; Addison's disease, neoplasm; tissue regeneration; chemotaxis; food additive; food preservative; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treating
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                                                                                                                             0.7%; Score 18.2; DB 1; Length 19;
94.7%; Pred. No. 8.6e+02;
ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hasel KW,
                                                                Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3' sequencing primer #1 used in the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glass J,
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                                                                                                                                                                                                                                                                                                                                   19 BAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2000; 2000US-0248892P.
28-NOV-2000; 2000US-0253623P.
                                                                                                                                                                     ilarity 94.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                        Local Similarity
es 18; Conserv
   present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD49149;
                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1033
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Matches
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                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Simplified TOGA method for simultaneous sequence-specific identification of multiple mRNA molecules in mRNA population, useful for determining tissue-specific patterns of gene expression or mechanisms of drug
                                                                                                                                                                                                                                                3' sequencing primer #1 used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                            Gaps
                                                                                                                                                                                                                                                                          drug interaction mechanism; drug screening; primer;
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14.7%; Pred. No. 8.6e+02;
ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chang HW;
 Length 19;
                            IndelB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hilbush BS, Hasel KWPD, Sutcliffe GJ,
0.7%; Score 18.2; DB 1; 94.7%; Pred. No. 8.6e+02;
                            ö
                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                 01-FEB-2002; 2002WO-US002666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94.78;
                                                                                                                                                               AAD50267 standard; DNA; 19
                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18; Conservative
                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-092784/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                Local Similarity
                                                                                                                                                                                                                                                                                 expression;
                                                                                                                                                                                                                                                                                                 genomic mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (QUAN/) QUAN J.
                                                                                                                                                                                                                                                                                                                                                         WO200261045-A2.
                                                                                                                                                                                                                                                                                                                            Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interaction.
                                                                                                                                                                                                                         24-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                      08-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Callahan MA;
                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19
                                                                                                                                                                                            AAD50267;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                        19
   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Quan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1035
ADC21495/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                 Matches
                                                                                                                                                                                                                                                                                   Gene
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Preparing a library of same-length signature sequences from a source nucleic acid population by ligating to the cleaved ends, a second adapter containing a recognition and cleavage site for a second restriction
                                                                                                              ss; tag-DNA signature; adapter-signature-adapter; parallel sequencing; genomic mapping; genetic identification; medical diagnostic.
                                                           DNA oligo (30) used in preparing a library of same length signatures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hiemisch H, Williams S,
Fu R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (LYNX-) LYNX THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                26-APR-2002; 2002US-0375782P.
                                                                                                                                                                                                                                                                                                                                                           25-APR-2003; 2003WO-US013076.
     (first entry)
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                                                                                                                                                                                                                                                  WO2003091416-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  endonuclease.
                                                                                                                                                                                             Unidentified
     26-FEB-2004
                                                                                                                                                                                                                                                                                                     06-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fischer A,
Vermaas E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention.
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Best Local S
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  à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel tumor-associated oligopeptide that is recognized as an antigen by CD8+ cytotoxic T lymphocytes (CTL) and causes CTL-induced lysis and/Or apoptosis of tumor cells, especially multiple myeloma cells. The oligopeptide is derived from human PRDI-BFI (Dositive regulatory domain I-binding factor-1) which is able to induce an MHC (major histocompatibility complex) class I allels variant A2-restricted immune response of CD8+ CTL against tumor cells. The products of the invention have cytostatic activity and can be used in a vaccine. The convention have cytostatic activity and can be used in a vaccine. The proping of the invention, also related retro-inverse and pseudopeptides, clusion proteins (FP), polymucleotides, vectors, host cells and antibodies and T cell receptors specific for PRDI-BFI peptides are useful for creating diseases associated with PRDI-BFI, particularly tumors. The products of the invention are also useful as diagnostic, therapeutic and corphylactic agents for detecting, modifying, generating, expanding activation and functional status of T cells, and for prophylactic agents for mono-clonal or recombinant A2-restricted T cell receptors and their functional equivalents. This sequence represents and for the products of poly- or mono-clonal or recombinant A2-restricted in the functional equivalents. This sequence represents and functional equivalents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                             tumor; antigen; CD8+ cytotoxic T lymphocyte; CTL; CTL-induced lysis; multiple myeloma cell; human; PRDI-BFI; positive regulatory domain I-binding factor-1; MHC; major histocompatibility complex Class I; cytostatic; vaccine; ss; primer; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New tumor-associated oligopeptide, useful particularly for treating multiple myeloma, is recognized by CD8 cytotoxic T cells, also derivatives and related nucleic acid.
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Pred. No. 8.6e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 22; 64pp; German.
BP.
                                                                                                                                                           Human PRDI-BF1 RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-SEP-2002; 2002WO-EP010701.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-SEP-2001; 2001DE-01048236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%;
ADC21495 standard; DNA; 19
                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IMMU-) IMMUGENICS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Theobald M, Lotz C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-354724/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003029282-A2
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                        18-DEC-2003
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Best Local Simi
Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-2003.
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                                                     ADC21495;
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Walker R;

Brenner S,

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                                                      This invention relates to a novel method for preparing a library of same-length signature sequences from a source nucleic acid population. Specifically, it comprises producing solid phase cloned libraries of oligonucleotide tag-DNA signature sequence constructs, which are useful for sequencing many polynucleotides simultaneously. The present invention constructs using first, and 'second' adapter-signature-adapter constructs using first, and 'second' adapters each containing a specific restriction endomuclease recognition site, and which flanks the same length signature sequence. As such, using the method described herein it is possible to do parallel sequencing of large populations of polynucleotides for genomic mapping, genetic identification and medical diagnostics. This oligonucleotide sequence is a DNA oligo involved in the step wase process of preparing a library of same length signature sequences from restriction fragments in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        infestinal epithelium cell development; peyer's patch M cell development; inflammatory bowel disease; glutenenteropathy; infectious disease; autoimmune disease; haemolytic anaemia; rheumatoid arthritis; dermatitis; Grave's disease; multiple sclerosis; allergy; asthma; diabetic mellitus; immune system disorder; hypersensitivity; anaphylaxis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Intestinal epithelium/peyer's patch M cell-related primer #15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ch 0.7%; Score 18.2; DB 1; Length 19; 1 Similarity 94.7%; Pred. No. 8.6e+02; 18; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
Disclosure, Fig 2a, 54pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2708 TAAAAAAAAAAAAAA 2726
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2708 TAAAAAAAAAAAAAAA 2726

19 BAAAAAAAAAAAAAA 1

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ADF74670 standard; DNA; 19 BP

RESULT 1036 ADF74670 ADF74670

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                                                                                                                                                                                                                                       The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the invention are also useful in the treatment of: inflammatory bowel disease, glutenenteropathy, infectious disease, autoimmume diseases (e.g. haemolytic anaemia, rheumatoid arthritis, dermatitis, Grave's diseases or discretes of the immumen system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequence represents a primer that was used in the exemplification of the
                                                                                                                                                                              Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function, useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RT-PCR primer used to amplify yeast clone-derived plasmid RNA
                                                                                                                                           2
                                                                                                                                                                                                                                                                                                                                                                                             0.7%; Score 18.2; DB 1; Length 19; 94.7%; Pred. No. 8.6e+02; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plant growth regulant, plant, agriculture; plant breeding;
transgenic plant; flowering; ss; RT-PCR; primer;
reverse transcriptase PCR.
                                                                                                                                          Mah SP,
                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                          O'mahony DJ, Evans CF,
 blood group incompatibility; ss; PCR; primer.
                                                                                                                                                                                                                        Disclosure; SEQ ID NO 360; 152pp; English.
                                                                                                                       (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                     2708 TAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                466/c
ADY39466 standard; DNA; 19 BP.
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                                                                               04-APR-2002; 2002WO-US010873.
                                                                                                   04-APR-2001; 2001US-0281416P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                           Byrne D,
                                                                                                                                                             WPI; 2003-075470/07.
                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CERE-) CERES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2005019462-A1.
                                         WO200280852-A2
                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAY-2005
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                                                                                                                                           Brayden DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADY39466;
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1038
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ADY39466/c
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The invention comprises the amino acid and coding sequences of lethal or non-viable proteins. The DNA and protein sequences of the invention are useful for transforming a plant cell, and producing transgenic plants that produce seeds that are not viable, not fertile, not capable of germinating, or are otherwise not capable of graminating, or are otherwise not capable of growing into mature plants. The present DNA sequence represents an oligo(dT)18 primer that was used
                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated nucleic acid molecule from Arabidopsis, Maize, or Brassica and encoding an amino acid sequence exhibiting at least 8% sequence identity to SEQ ID NO. 3, 5, 7, 10, 12, 14, 17, 19, 21, 24, 26, 28, 31, 34, 36, 38, 41, 43, 45, 48, or 49 as given in the specification. The nucleic acids and polypeptides of the invention may act as plant growth regulators and as such may be useful for generating transgenic plants having increased size, increased number and size of tosette leaves and are late-flowering. The current sequence is that of the RT-PCR primer of the invention which was used to amplify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel polypeptide having characteristic of being lethal or non-viability polypeptide, useful for producing transgenic plants that produce seeds that are not viable, not fertile, and are not capable of germinating.
                                                                                                                       Brassica, useful for generating transgenic plants having increased size, increased number and size of rosette leaves and are late-flowering.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Non-viable seed-producing transgenic plant-related-oligo(dT)18 primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transgenic plant; seed; artificial seed; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
   Zhang H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18.2; DB 1;
94.7%; Pred. No. 8.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
   Dang V,
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                                                                                                                                                                                                                                                           Disclosure; Page 63; 151pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                yeast clone-derived plasmid RNA.
   Kwok S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19 BAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 94.7
es 18; Conservative
Pennell R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Feldman K, Nadzan G,
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                                                                  WPI; 2005-214253/22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2005035763-A1.
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         Feldman K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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Query Match

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RESULT 1040

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The invention relates to a nucleic acid molecule or its complement sequence capable of modulating transcription. The nucleic acid molecule of the invention is useful for transcription of polynucleotides in a host cell or transformed host organism. The present sequence is a primer used for modulating gene transcription.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated nucleic acid molecule capable of modulating transcription, or its complement, useful for transcription of polynucleotides in a host cell or transformed host organism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primer d(T)18, SEQ ID NO: 73, used to generate probes for hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transgenic plant; plant growth regulation; development; food,
agriculture; horticulture; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.7%; Score 18.2; DB 1; Length 19; 94.7%; Pred. No. 8.6e+02; vative 1; Mismatches 0; Indels
                                                                                                   Oligo(dT)18 primer used for modulating gene transcription.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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Pennell R, Schneeberger R, Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, SEQ ID NO 1; 210pp; English.
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06-NOV-2003; 2003US-0518075P.
04-DEC-2003; 2003US-05293511P.
12-DEC-2003; 2003US-0529352P.
13-FEB-2004; 2004US-05444771P.
30-JUN-2004; 2004US-0583691P.
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                                                                                                                                                                                                                                                                                                                               2003US-0505689P
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                                                                (first entry)
                                                                                                                                        Transcription; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-664198/68.
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                                                                01-DEC-2005
                                                                                                                                                                                                                                                                                                                               23-SEP-2003;
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                                                                                                                                                                                                                                                     06-OCT-2005
                             AED19813;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated nucleic acid promoter or promoter control element sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AEC21692) which are capable of modulating transcription. The sequences are useful as promoter sequences or promoter control element sequences for modulating transcription, e.g. constitutive transcription, stress induced transcription, light induced transcription, dark induced transcription, leaf transcription, root transcription, silique transcription, callus transcription, flower transcription, immature bud and inflorescence specific transcription, or senescing induced transcription. The present sequence is a primer used for generating probes for the promoter sequences of the invention.
                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               useful for modulating transcription, e.g. constitutive transcription, stress induced transcription, light induced transcription, or callus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to novel promoter sequences (AEC21689-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                          Score 18.2; DB 1; Length 19;
Pred. No. 8.6e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.7%; Score 18.2; DB 1; Length 19; Best Local Similarity 94.7%; Pred. No. 8.6e+02; Matches 18; Conservative 1; Mismatches 0; Indels
                                     Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Feldmann K;
in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 1; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Promoter; transcription; primer; ss.
                                                                                                                                                  2708 TAAAAAAAAAAAAAA 2726
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                                                                                                                                                                       Medrano L,
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AEC21688 standard; DNA; 19 BP.
                                                                          0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-OCT-2003; 2003US-0511460P
                                                                                                                                                                                                                                                                                                                                                                                                 Oligo dTV primer, SEQ ID 1
                                                                                                                                                                                                                                                                                                                                                          17-NOV-2005 (first entry)
                                                                                             Best Local Similarity 94.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Penell R, Apuya N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2005-618221/63
                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transcription.
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Medrano L;

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Kwok S,

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06-OCT-2005

AED19813 standard; DNA; 19 BP.

RESULT 1041 AED19813/c ID AED190

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The invention relates to an isolated nucleic acid molecule capable of modulating transcription, where the nucleic acid molecule shows at least modulating transcription, where the promoter sequences given in the specification or its complement. The invention also relates to a vector construct comprising a first nucleic acid capable of modulating transcription, where the nucleic acid molecule shows at least 80% transcription, and a second nucleic acid molecule shows at least 80% sequence identity to one of the promoter sequences given in the specification, and a second nucleic acid molecules are heterologous to each the first and second nucleic acid molecules are heterologous to each contract and are operably linked together, a host cell comprising the other and are operably linked together, a host cell comprising the comprising the vector construct. The first conception and a plant comprising the vector construct. The first conception and a plant comprising the vector construct. The first conception and a given in the specification, where the first nucleic acid molecule is inserted into a plant cell and the plant cell is regenerated concerned and promoter control elements are useful for modulating transcription of a desired polynucleotide in plants. This sequence represents a synthetic continer used in the scope of the invention.
      Disclosure; SEQ ID NO 1; 787pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1044
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                  The present invention relates to polynucleotides and their encoding polypeptides with the use of those products for making transgenic plants. The sequences of the invention are useful modifying plant characteristics or for modulating and manipulating growth, development and biochemistry of a plant. The invention is useful for producing plants with increased yield of biomass or chemical components, in particular food and reproducible raw materials. The present sequence is a d(T)18 primer used to generate probes for hybridization. This sequence is used in making
                                                                                                                                                                                   for
                                                                                                                                                                              New nucleotide seguences, useful modifying plant characteristics or for modulating and manipulating growth, development, and biochemistry of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                        Kwok S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.7%; Score 18.2; DB 1; Length 19; Best Local Similarity 94.7%; Pred. No. 8.6e+02; Matches 18; Conservative 1; Mismatches 0; Indels
                                                                                                         Kiegle EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ľu Y,
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                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                           Feldmann K,
Wu C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cook Z, Fang Y, Feldmann K, Kiegle EA,
Pennell R, Schneeberger R, Wu C;
                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 73; 132pp; English.
                                                                                                           Alexandrov N, Zhihong C, Fang Y,
Lu Y, Penell R, Schneeberger R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transcription; vector; primer; ss.
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04-DEC-2003; 2003US-0527611P.
13-FEB-2004; 2004US-0544771P.
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                     23-SEP-2004; 2004US-00950095
                                                   23-SEP-2003; 2003US-0505420P
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                                                                                                                                                        WPI; 2005-683371/70.
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                                                                                    (CERE-) CERES INC.
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                                                                                                                                                                                                                        plant.
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                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated nucleic acid molecule ({f I}) ,
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                                                Query Match 0.7%; Score 18.2; DB 1; Length 19; Best Local Similarity 94.7%; Pred. No. 8.6e+02; Matches 18; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plant; transgenic plant; herbicide resistance; ss; primer.
Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 1; 157pp; English.
                                                                                                                                                                 2708 TAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plant promoter associated primer.
                                                                                                                                                                                                                      BAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-2004; 2004US-0564658P.
23-APR-2004; 2004US-0564678P.
                                                                                                                                                                                                                                                                                                                                                                 AED87374 standard; DNA; 19
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19-JAN-2006.
                                                                             Kiegle EA,
                                                         Synthetic.
                                            AEF26613;
                        Query Match
                                      RESULT 1045
                           Matches
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Query Match 0.7%; Score 18.2; DB 1; Length 20; Best Local Similarity 94.7%; Pred. No. 8.9e+02; Matches 18; Conservative 1; Mismatches 0; Indels ö comprising a plant promoter sequence. (I) is useful for producing a transformed plant having shade responsive characteristics different from those of a naturally occurring plant of the same species cultivated under the same conditions, which involves introducing (I) into a plant or plant cell to modulate transcription in a plant. (I) is useful for expressing a developmental DNA sequence in a plant. (I) is useful for understanding developmental mechanisms e.g. shade responsive promoters that are induced during callus formation and somatic embryo formation, and for isolating trans-acting factors. (I) is also useful for modulating transcription in modulating genes involved in defense, pest resistance, herbicide resistance etc. The present sequence represents a plant promoter associated primer. The present invention relates to isolated nucleic acid molecules from Arabidopsis thaliana and the polypeptides encoded by them. The patentees also claim a vector construct containing such nucleic acid molecules, and a host cell transformed; a method for detecting a nucleic acid in a sample; and a plant, plant cell, plant material or seed of a plant which comprises the nucleic acid molecule which is exogenous or heterologous to the plant or plant cell. The vector construct comprises a first nucleic acid having a regulatory sequence capable of causing transcription and/or translation in a plant and a second nucleic acid having the sequence of the isolated nucleic acid molecule. The first and second nucleic acids are operably linked and where the second nucleic acid is heterologous to any element in the vector construct. The isolated nucleic acid molecules are useful in producing transgenic plants for use as models for modifying New nucleic acid molecule, useful in producing transgenic plants for use as models for modifying plant characteristics e.g. increase in plant height, number or size of leaves, or wood products. Gaps Jofuku D; . 0 / Match 0.7%; Score 18.2; DB 1; Length 19; Local Similarity 94.7%; Pred. No. 8.6e+02; nes 18; Conservative 1; Mismatches 0; Indels Schneeberger R, Fang Y, Kwok S, Apuya N; Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other; Example 1; SEQ ID NO 132; 245pp; English. 2708 TAAAAAAAAAAAAAA 2726 AEF26613 standard; DNA; 19 BP 09-DEC-2004; 2004US-00010239. 88; primer; transgenic plant. 12-DEC-2003; 2003US-0529352P. 23-MAR-2006 (first entry) Pennell R, Okamuro J, Donson J, WPI; 2006-099536/10. Oligo(dT)18 primer (CERS-) CERS INC. US2006015970-A1.

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This invention descibes a novel method for the analysis of a DNA fragment which comprises: (i) addition of a known common oligonucleotide sequence to at least one terminal of each DNA fragment, (ii) amplification of the resultant DNA fragment as a primer using a first common primer containing a complementary nucleotide sequence to the above mentioned known common of oligonucleotide sequence to the prepared known common componed to the sequence optionally having been introduced with complementary nucleotide sequence at a terminal, and a specific primer companies of the prepared known common oligonucleotide sequence to the prepared known common componed to the properties of the properties of the amplified DNA to find the information of the DNA fragment, in which the specific primer is designed to prepare fragments of the common first and second primers and to give short fragment of amplified DNA and (iv) contrament to make their differentiation. Of the DNA fragment of amplified DNA and of the common first and second primers and to give short fragment of amplified DNA and (iv) confirmations of known and unknown genes readily provides information of unknown genes and simultaneous monitoring of signals derived from minor confirmation of the properties of known and unknown according to functions of known and unknown according to functions of known and confirmation of signals derived from minor confirmation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis of DNA fragment - comprises addition of known common oligonucleotide, amplification of resultant DNA fragment and analysis and labelling of amplified DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes can be performed. AA209189-Z09201 represent oligonucleotide primers
plant characteristics, e.g. increase in plant height, number or size of leaves, or wood products. The present sequence is that of a primer used in generation of labeled probes from first-stand cDNA.
                                                                                                                                                                                Gaps
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                                                                                                                             Ouery Match 0.7%; Score 18.2; DB 1; Length 19; Best Local Similarity 94.7%; Pred. No. 8.6e+02; Matches 18; Conservative 1; Mismatches 0; Indels
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                                                                                   Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide 9 for DNA analysis.
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                                                                                                                                                                                                                     2708 TAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                 AAZ09197 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Solid phase immunoassay using oligo:nucleotide as label - also new conjugates of oligo:nucleotide coupled to antigenic peptide, partic. for diagnosing hepatitis C or E virus infection.
                                                                                                                    Synthetic oligo; solid phase immunoassay; ss
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                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                        Khudyakov YE;
                               (revised)
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-006819/01.
                                                                                                                                                                                                                                                                 13-MAY-1994;
                                                                                                                                                                                                                                                                                                    13-MAY-1993;
                                                                                                                                                                                             WO9426932-A1
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                               25-MAR-2003
03-AUG-1995
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                                                                                      PCR primer
                                                                                                                                                          Synthetic.
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 AAQ75025;
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consecuing a library of bovine Mbol DNA fragments of between 250 and 500

consecuent an (AC)15 and a (TC)15 oligonuclectide probe. One out of 50

clones cross-hybridised. Assuming independent distribution of

microsatellites and Mbol sites, the frequency of (T6)n >9 microsatellites

consecuent and proper second of the frequency of (T6)n >9 microsatellites

consecuent and indexed herein (see below). The sequence information

for ca. 230 such bovine microsatellites is summarised in the

specification and indexed herein (see below). The sequences upstream and

comparream of the microsatellite sequence were used to generate the

required PCR primers for in vitro amplification of the corresp.

microsatellite (using the program OPTIPRIM). The microsatellites may be

microsatellite (using the program OPTIPRIM). The microsatellites may be

consecuently important trait loci, or genes involved the determinism of

economically important traits esp. in cattle, to allow selective

breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                      PCR; selection; primers, OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / Match 0.7%; Score 18; DB 1; Length 18; Local Similarity 100.0%; Pred. No. 8.7e+02; les 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                     Sequence of a microsatellite from clone TGLA60B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Table 7; Page 375; 517pp; English.
                       2708 TAAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВАААААААААААААА 1
                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92WO-US000340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91US-00642342
                                                                                                                                                   AAQ34110 standard; DNA; 18
                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 маввеу ЛМ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENM-) GENMARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LS-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                    WO9213102-A1
                                                                                                                                                                                                                     25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                       06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Georges M,
                                                                                                                                                                                                                                                                                                                                                                   Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                      AAQ34110;
                                                                                                                RESULT 1047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         datches
                                                                                                                                    AAQ34110
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AAR62941 and AAR62942 are examples of synthetic immunoreactive peptides.

CT They are used in a method for detecting an antigen in a subject. The method involves binding the antigen to a solid support and then reacting it with an immunoreactive ligand (L) bound to an oligo; removing any curreacted L, and then detecting the presence of the oligo. A similar curreacted L, and then detecting the presence of the oligo. A similar captured and by the see the ligand is an oligo. The use of an amplifiable oligo as the label allows Ag or be labelled Ag. The use of an amplifiable oligo as the label allows Ag or to be detected at very low levels. An exemplary olgi is AAQ75024 which can be covalently attached by the 5· termina to the N· or C·terminal of can synthetic peptide. In the example, peptide AR62941 was coupled to oligo AAQ75024 using disuccinimidyl suberate. Serum samples suspected to contain HBV Abs were immobilised on plastic tubes or wells, then contain HBV Abs were immobilised on plastic tubes or wells, then captide-oligo product. The vessels were separate tube using as primers AAQ75025 and AAQ75026 in 30 cycles of PCR. The amplification product - AAQ75031 - was treated with uracil DNA cycles of product captured by immoblised oligo-dT. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Flavonoid 3' hydroxylase; pigmentation; flower colour; transgenic plant; snapdragon; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 0 A; 0 C; 0 G; 0 T; 18 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Score 18; DB 1; Le
100.0%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anchored poly(T) oligonucleotide polyT-AnchA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT94667 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT94667;
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AAQ75025 standard; RNA; 18 BP.

RESULT 1048

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AAQ75025,

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Sequence 18 BP; 1 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                              (OLIG-) OLIGOS ETC & OLIGOS THERAPEUTICS INC.
                                                                                Michael MZ;
                                                                                                                          Example 15; Page 59; 234pp; English.
                                                                                                                                                                                                                                          2708 TAAAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                          Thompson TL;
                                                                                                                                                                                                                                                                                     AAV21970 standard; DNA; 18 BP
                                          97WO-AU000124.
                                                      96AU-00008386.
                                                                                                                                                                                                                                                                                                                                                                                                     97WO-US012961
                                                                                                                                                                                                                                                                                                                                                                                                                  96US-00685575
                                                                                                                                                                                                                                                  14-JUL-1998 (first entry)
                                                                               Holton TA,
                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 18; Conservative
                                                                  (FLOR-) FLORIGENE LTD.
                                                                                            WPI; 1997-448691/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Dale RMK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-120687/11
                                                                               Brugliera F,
                                                      01-MAR-1996;
                 WO9732023-A1
                                          28-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                     23-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                  24-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                            WO9803533-A1
                              04-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-1998
     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                 AAV21970;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Arrow A,
                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                         RESULT 1050
                                                                                                                                                                                                                                                                               AAV21970,
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                                                                                                                                                                                                                                                     셤
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Treating bacterial infections in humans or animals with (KAGA ) ZH KAGAKU & KESSEI RYOHO KENKYUSHO Primer SEQ ID NO:3 from JP11075880. Claim 49; Page 87; 163pp; English. 2709 AAAAAAAAAAAAAAA 2726 18 AAAAAAAAAAAAAA 1 98JP-00195719. 97JP-00205378. AAX19943 standard; DNA; 18 (first entry) Conservative enhances cellular uptake Query Match Best Local Similarity Matches 18; Conservat WPI; 1999-257710/22 JP11075880-A. 10-JUL-1998; 14-JUL-1997; 14-JUN-1999 23-MAR-1999. Synthetic. AAX19943; RESULT 1051 AAX19943 ò 요 ö Anchored poly(T) oligonucleotides polyT-anchA (AAT94667), polyT-anchC (AAT94668) and polyT-anchG (AAT94669) are complementary to the upstream region of a polyadenylation sequence. They were used to prime cDNA synthesis from snapdragon (Antirrhinum majus) petal and leaf RNA, and were also utilised in the PCR amplification of plant cytochrome P450 sequences (see also AAT9470-73). A CDNA clone (see AAT94657) encoding flavonoid 3' hydroxylase (see AM35704) was isolated using a differential display approach. This can be used to manipulate the pigmentation of transgenic plants Novel flavonoid 3'-hydroxylase(s) from flowering plants - and corresponding DNA, used in the manipulation of pigmentation in plants. Gaps Nuclease resistant antisense oligo NBT 13 targeted against (T)18. ö Nuclease resistant; bacterial infection; antibiotic; target; veterinary medicine; treatment; human; industrial process; bacterial control; ss. 0.7%; Score 18; DB 1; Length 18; 00.0%; Pred. No. 8.7e+02; 0; Indels 100.0%; Prec. ...

The treatment of animals, including humans, having a bacterial infection.

The treatment comprises administration of such nuclease resistant
oligonucleotides, targeted to a nucleic acid or protein of the bacterium,
and formulated with a carrier. A compound comprising this nuclease
resistant oligonucleotide can be covalently linked to an antibiotic. The
resistant oligonucleotide can be covalently linked to an antibiotic. The
method is used to treat infections by a wide variety of Gram-positive and
Gram-negative, or acid-fast, bacteria, in human and veterinary medicine.
The methods are particularly used in immuno-compromised individuals (e.g.
The methods are particularly used in immuno-compromised individuals (e.g.
The methods are particularly used in immuno-compromised individuals (e.g.
The methods are particularly used in immuno-compromised individuals (e.g.
The methods are particularly used in immuno-compromised individuals (e.g.
The methods are squired immunofeficiency syndrome or those receiving
Chemotherapy or radiation therapy), optionally in combination with, or
fused to, antiviral or other antimicrobial oligonucleotides. Apart from
Chaboratory cultures, foods, beverages and industrial processes. The
oligonucleotides are specific for bacteria, without affecting metabolism
on mammalian cells. They may also activate Rhase H and have a general,
non-specific immune-stimulating effect. The oligonucleotides can be
administered orally, intranasally, rectally, topically or by injection,
optionally coupled to an agent (e.g. carbohydrate or polyamine) that ö This antisense oligonucleotide is nuclease resistant and can be used in the treatment of animals, including humans, having a bacterial infection oligo:nucleotide(s) - resistant to nuclease and targetted to bacterial nucleic acid or proteins, also conjugates of these oligo:nucleotide(s) with antibiotics. A method has been developed for labelling an oligonucleotide having a repeated sequence of (XY)n (where X and Y consists of a combination of adenine and thymine or uracil or guanine and cytosine, and n is an integer of 1 or more ) at the 3'-terminal side in which the repeated Gaps Primer; oligonucleotide; labelling; detection; self-priming; PCR; ss. ö Labelling of an oligonucleotide - useful for detecting genes. 0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.78+02; tive 0; Mismatches 0; Indels Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other; Example 1; Page 7; 10pp; Japanese. 

ngs.res

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Tue Nov

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Beta-D-arabinose, antisense; inhibition; transcription; expression; reverse transcription; viral replication; RNase H cleavage; triple helix formation; ss.

Oligoarabinonucleotide SEQ ID NO:2

(first entry)

08-MAY-2000

AAZ87161;

a "Ribose moiety replaced by beta-D-arabinose"

Location/Qualifiers

1. .18 /\*tag= /note= "

Key modified\_base

Synthetic

WO9967378-A1

29-DEC-1999.

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ö
                    constituting the repeated sequence and a DNA polymerase lacked in 5' to 3' exonuclease activity. The method can be used for detecting a gene. The method can detect a gene in a sensitivity up to ten times higher than prior art methods. The present sequence represents a primer used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A method has been developed for labelling an oligonucleotide having a repeated sequence of (XY)n (where X and Y consists of a combination of adenine and thymine or uracil or guanine and cytosine, and n is an integer of 1 or more ) at the 3'-terminal side in which the repeated sequence is added and extended using a labelled body of the nucleotide constituting the repeated sequence and a DNA polymerase lacked in 5' to 3' exonuclease activity. The method can be used for detecting a gene. The method can detect a gene in a sensitivity up to ten times higher than prior art methods. The present sequence represents a primer used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prior art methods. The present sequence represents a primer used in an example from the present invention
sequence is added and extended using a labelled body of the nucleotide
                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer; oligonucleotide; labelling; detection; self-priming; PCR; ss.
                                                                                                                                                                                                                   ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Labelling of an oligonucleotide - useful for detecting genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 18; DB 1; Length 18; 00.0%; Pred. No. 8.7e+02;
                                                                                                                                                                            DB 1; Length 18;
8.7e+02;
                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                      Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Pred. ....
                                                                                                                                                                        Query Match 0.7%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 8.7 Matches 18; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KAGA ) ZH KAGAKU & KESSEI RYOHO KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer SEQ ID NO:2 from JP11075880.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 7; 10pp; Japanese
                                                                                                                                                                                                                                                           2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                             AAX19942 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98JP-00195719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97JP-00205378
                                                                                                                                                                                                                                                                                               18 AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-257710/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP11075880-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX19942;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                         RESULT 1052
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Therapeutic composition containing antisense oligonucleotides that include arabinose sugars, particularly for inhibiting viral replication.

Example 1; Page 29; 91pp; English.

Arion D;

Borkow G,

Wilds C,

Noronha AM,

Damha MJ, Parniak MA, (UYMC-) UNIV MCGILL.

WPI; 2000-160584/14.

99WO-CA000571 98CA-02241361

17-JUN-1999; 19-JUN-1998;

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The invention relates to a new composition for selective, sequence-
ce specific inhibition of gene transcription and expression in a host. The
composition comprises oligonucleotides containing arabinose sugars that
can hybridise to either a a single-stranded (ss) RNA to induce RNase H
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
ce lelix, thereby inhibiting DNA replication and/or transcription. The
coligoarabinonucleotides are used for antisense inhibition of gene
cypression or to prevent DNA replication, or reverse transcription of
control prevent DNA replication, or reverse transcription of
control in combination with RNase H, as reagents for sequence-specific
cinhibit retrovirus replication. The oligoarabinonucleotides can also be
consequent to make mapping, and additionally for the study and control of
central genomic RNA, increased resistance to nucleases and show little if
affinity for RNA, increased resistance to nucleases and show little if
cany non-specific binding to cellular or serum proteins They target as
controlleotides containing pyrimidine bases form triple helices
coligoarabinonucleotides containing pyrimidine bases form triple helices
coligoarabinonucleotides containing pyrimidine bases form triple helices
coligoarabinonucleotides containing bear-barabinese used in an
colligoarabinonucleotides containing bear-b-arabinose used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
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RESULT 1054 AAZ87162/c

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Gaps ; 0

0; Indels

2709 AAAAAAAAAAAAAA 2726

Best Loca Matches

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AAZ87161 standard; RNA; 18 BP.

RESULT 1053

AAZ87161 ID AAZR

2'-deoxy-2'-fluoro-beta-D-arabinose; antisense; inhibition; transcription; expression; reverse transcription; viral replication; RNase H cleavage; triple helix formation; ss.

Deoxyarabinonucleotide SEQ ID NO:7.

(first entry)

08-MAY-2000

AAZ87166;

AAZ87166 standard; DNA; 18 BP.

AAZ87166/c

.18 //rtag= a /notce= "Deoxyribose moiety replaced by 2'-deoxy-2'-fluoro-beta-D-arabinose"

Location/Qualifiers

modified\_base

Synthetic

WO9967378-A1.

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Tue Nov

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Therapeutic composition containing antisense oligonucleotides that include arabinose sugars, particularly for inhibiting viral replication
                                                      Beta-D-arabinose; antisense; inhibition; transcription; expression; reverse transcription; viral replication; RNase H cleavage; triple helix formation; ss.

    .18
    /*tag= a
    /note= "Ribose moiety replaced by beta-D-arabinose"

                                                                                                                                                                                                       Arion D;
                                                                                                                                                                                                       Borkow G,
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 0 A; 0 C; 0 G; 0 T; 18 U; 0 Other;
                                                                                                                                                                                                      Noronha AM, Wilds C,
                                                                                                                                                                                                                                                                                                                                                                                                                  exemplification of the present invention
                                                                                                Location/Qualifiers
                                         Oligoarabinonucleotide SEQ ID NO:3.
                                                                                                                                                                                                                                                     Example 1; Page 29; 91pp; English.
 BP.
                                                                                                                                                             99WO-CA000571
                                                                                                                                                                          98CA-02241361.
AAZ87162 standard; RNA; 18
                           (first entry)
                                                                                                                                                                                                      Parniak MA,
                                                                                                                                                                                       (UYMC-) UNIV MCGILL.
                                                                                                                                                                                                                   WPI; 2000-160584/14.
                                                                                               Key
modified_base
                           08-MAY-2000
                                                                                                                                                                         19-JUN-1998;
                                                                                                                                 WO9967378-A1
                                                                                                                                                            17-JUN-1999;
                                                                                                                                               29-DEC-1999.
                                                                                  Synthetic
                                                                                                                                                                                                      Damha MJ,
             AAZ87162;
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Therapeutic composition containing antisense oligonucleotides that include arabinose sugars, particularly for inhibiting viral replication.

Example 2; Page 31; 91pp; English.

Arion D;

Borkow G,

Wilds C,

Noronha AM,

Parniak MA,

Damha MJ,

WPI; 2000-160584/14.

(UYMC-) UNIV MCGILL.

99WO-CA000571. 98CA-02241361.

17-JUN-1999;

19-JUN-1998;

```
The invention relates to a new composition for selective, sequence-
specific inhibition of gene transcription and expression in a host. The
composition comprises oligonucleotides containing arabinose sugars that
can hybridise to either a a single-stranded (ss) RNA to induce RNase H
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage or the oreompositions are therefore particularly used to
clinhibit retroviral replication. The oligoarabinonucleotides can also be
cleavage or RNA mapping, and additionally for the study and control of
cleavage or RNA mapping, and additionally for the study and control of
cgene expression in cells. The oligoarabinonicleotides have excellent
affinity for RNA, increased resistance to nucleases and show little if
can non-specific binding to cellular or serum proteins. They target ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA, but not complementary so DNA, so may be useful for targetting retroviral genomic RNA to inhibit the early stages of viral replication. Oligoarabinomucleotides containing pyrimidine bases form triple helices with significantly higher thermal stability than those produced by normal oligoancleotides. Sequences AAZ87165-287169 represent oligodeoxyarabinomucleotides containing 2'-deoxy-2'fluoro-beta-barabinose used in an exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 18; DB 1; Le
100.0%; Pred. No. 8.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
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Gaps

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0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; ive 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAAA 2726

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Best Local Similarity Matches 18; Conserv

Query Match

8

ngs.res

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The invention relates to a new composition for selective, sequence-
specific inhibition of gene transcription and expression in a host. The
composition comprises oligonuclectides containing arabinose sugars that
can hybridise to either a a single-stranded (ss) RNA to induce RNase H
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
cleavage activity, or to a DNA/DNA or DNA/RNA duplex to form a triple
oligoarabinonuclectides are used for antisense inhibition of gene
expression or to prevent DNA replication, or reverse transcription. The
by retroviruses. The compositions are therefore particularly used to
inhibit retroviral replication. The oligoarabinonucleotides can also be
used, in combination with RNase H, as reagents for sequence-specific
cleavage or RNA mapping, and additionally for the study and control of
gene expression in cells. The oligoarabinonucleotides have excellent
cf gene expression in cells. The oligoarabinonucleotides have excellent
caffinity for RNA, increased resistance to nucleases and show little if
any non-specific binding to cellular or serum proteins. They target as
RNA, but not complementary as DNA, so may be useful for targetting
cretroviral genomic RNA to inhibit the early stages of viral replication.
Oligoarabinonucleotides containing pyrimidine bases form triple helices
with significantly higher thermal stability than those produced by normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Therapeutic composition containing antisense oligonucleotides that include arabinose sugars, particularly for inhibiting viral replication.
                                                                                                                                                                                       2'-deoxy-2'-fluoro-beta-D-arabinose; antisense; inhibition;
transcription; expression; reverse transcription; viral replication;
RNase H cleavage; triple helix formation; ss.
                                                                                                                                                                                                                                                                                                                                                /*tag= a
/note= "Deoxyribose moiety replaced by 2'-deoxy-2'-
fluoro-beta-D-arabinose"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arion D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2'-deoxy-2'fluoro-beta-D-
the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Borkow G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Noronha AM, Wilds C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligodeoxyarabinonucleotides containing arabinose used in an exemplification of
                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                    Deoxyarabinonucleotide SEQ ID NO:8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 31; 91pp; English.
                                    AAZ87167 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-CA000571.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98CA-02241361.
                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Damha MJ, Parniak MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-160584/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYMC-) UNIV MCGILL.
                                                                                                                                                                                                                                                                                                                  Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUN-1998;
                                                                                                                  08-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9967378-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-DEC-1999
                                                                                                                                                                                                                                                                           Synthetic.
                                                                           AAZ87167;
RESULT 1056
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Oligonucleotide #6 used for the preparation of normalised cDNA libraries.
                                                                                                                                                                         Rat; secreted factor; clone P0018B D12; cardiant; antiinflammatory; antiarrhythmic; antiarteriosclerotīc; antiatherosclerotic; nephropathic; antidiabetic; immunosuppressive; antiasthmatic; antisheumatoid; antidiabetic; immunosuppressive; antiasthmatic; antianterial; osteopathic; cerebroprotective; vasotropic; antiulcer; nootropic; neuroprotective; congestive heart failure; myocardial infarction; hypertrophic cardiomyopathy; angina pectoris; myocardial infarction; kidney disease; acute renal failure; renal glucosuria; renal infarction; polycystic kidney disease; hereditary nephritis; inflammatory disease; tumour angiogenesis; osteoarthritis; toxic shock syndrome; psorlasis; stroke; neural trauma; cerebral malaria; Crohn's disease; osteoporosis; ulcerative colitis; Alzheimer's disease; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel secreted factor encoded by clone P00188D12 which is differentially expressed in certain disease states, useful in diagnosing and treating cardiac, renal or inflammatory diseases.
1 AAAAAAAAAAAAAA 18
                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                          27-SEP-2000; 2000WO-US026544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0156280P.
                                                                   AAD03565 standard; DNA; 18
                                                                                                                           19-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stanton LW, Kapoun AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-266159/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SCIO-) SCIOS INC.
                                                                                                                                                                                                                                                                                                                                                         Rattus norvegicus.
                                                                                                                                                                                                                                                                                                                                                                                   WO200123564-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                              05-APR-2001.
                                                                                                 AAD03565;
                                          RESULT 1057
                                                         AAD03565
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DB 1; Length 18; 8.7e+02; Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other; Score 18; Pred. No. Query Match 0.7%; Sest Local Similarity 100.0%;

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Gaps

; 0

0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; ive 0; Mismatches 0; Indels

Query Match
Best Local Similarity 100.
Matches 18; Conservative

rat secreted factor P00188\_D12 gene

The patent discloses novel secreted factor protein encoded by clone pools 112. The secreted factor is differentially expressed in certain disease states. Secreted protein, its antibodies, antagonists or compositions comprising them are useful in the diagnosis and treatment of cardiac diseases such as congestive heart failure, myocardiis, hypertrophic cardiomyopathy, angina pectoris, myocardiis, and treatisocalerosis, widney diseases such as acute renal cardium, renal infarction, nephrogenic diabetes insipidus, polycystic kidney disease, hereditary nephritis and inflammatory diseases such as asthma, autoimmune diabetes, tumour angiogenesis, rheumatoid arthritis, osteoarthritis, toxic shock syndrome, asthma, stroke, neural trauma, psoriasis, cerebral malaria, osteoporosis, crohn's disease, ulcerative colitis, Alaheimer's disease. Secreted crohn's disease, ulcerative colitis, Alaheimer's disease. Secreted to reference RNA or DNA encoding the secreted factor is useful for detecting cardiac, kidney and inflammatory disease. The present DNA sequence is an oligonucleotide which is used in the preparation of a commalised convalined are used in the identification of differentially expressed convalined are used in the identification of differentially expressed

Example 1; Page 42; 71pp; English.

2709 AAAAAAAAAAAAAAA 2726 ||||||||||||||||

Matches

AAD17014;

AAD17014

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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
                                                                                                                                                                                       Vaccine, cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; tungal infection; parasitic infection; cancer; asthma; infections disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 18; DB 1; Length 18;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. wc.
                                                                                                                                                Immunostimulatory nucleic acid #824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schetter C, Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAAAA 2726
AAF99708/c
ID AAF99708 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18 AAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0156113P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                                                                                                                                                                                                                          25-SEP-2000; 2000WO-US026383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (IOWA ) UNIV IOWA RES FOUND. (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%;
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                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-273485/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                        WO200122972-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-1999;
27-SEP-1999;
                                                                                                       12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                 05-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Krieg AM,
                                                                                                                                                                                                                                                                                               Synthetic.
                                                                AAF99708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF99734;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF99734/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to an array of proteins (antibody mimics) comprising a fibronectin type III domain having a randomised loop, a randomised beta-sheet, or their combination, and has the capacity to bind to a compound that is not bound by a corresponding naturally-occurring fibronectin, immobilised onto a solid support. The antibody mimics is useful for detecting a compound preferably a protein, in a biological sample. It is also useful to detect one or more different analytes simultaneously in a sample. Hence is useful for diagnostic purposes. It is also useful for the purpose of designing proteins capable of binding to virtually any compound of interest. The present sequence is an oligonucleotide A18-2PEG linker used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fibronectin scaffold protein array for obtaining a protein/compound which binds to a compound/protein, comprises a fibronectin type III domain having a randomized loop, a randomized beta-sheet or their combination.
    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
    ;
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                                                                                                                                                                                                                                                                                                                                                       Scaffold protein; antibody mimic; fibronectin type III domain; randomised loop; randomised beta-sheet; diagnostic purpose;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0; Indels
    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag= a
/note= "Linked to (PEG)2CCPuromycin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
    ;
0
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kuimelis RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 25; 67pp; English.
                                           2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                Oligonucleotide A18-2PEG linker.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2001; 2001WO-US006414.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-FEB-2000; 2000US-00515260
                                                                                                                                                                                     AAD17014 standard; DNA; 18
                                                                                                                                                                                                                                                                     (first entry)
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lipovsek D, Wagner RW,
                                                                                                                                                                                                                                                                                                                                                                             randomised loop; rando
protein designing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-557782/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHYL-) PHYLOS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200164942-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        misc feature
                                                                                                                                                                                                                                                                       29-NOV-2001
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Gaps

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0; Indels

Immunostimulatory nucleic acid #850.

Query Match

Best Loca Matches

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RESULT 1059

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The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrinidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bantigens antigens antigens antigens. The method is staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Thi immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
             Vaccine, cytostatic, virucidal, bactericidal, fungicidal, anti-parasitic, immunostimulatory, tumour; viral infection; bacterial infection; fungal infection; parasitic infection; asthma; almin infection allergy, immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Phagemid vector; pCR2.1; rat; secreted factor; P00210D09; cardiant; nephrotropic; antiinflammatory; gene therapy; cardiac disease; renal disease; inflammatory disease; polylinker; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phagemid vector pCR2.1 polylinker oligonucleotide #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                  Krieg AM, Schetter C, Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.7%; Scc
Local Similarity 100.0%; Pr
tes 18; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF82472 standard; DNA; 18 BP
                                                                                                                                                                                                                                                    25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                  25-SEP-2000; 2000WO-US026383.
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                                                                                                                                                                                                                                                                                                                                               (COLE-) COLEY PHARM GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-273485/28
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                                                                                                                                             WO200122972-A2
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                                                                                                                                                                                05-APR-2001
                                                                                                           Synthetic.
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PUREL 1. It was used in the construction of a normalised rat cDNA library, which was used in an example demonstrating differential expression of a rat gene referred to as clone P00210009. The invention relates to a polypeptide comprising a sequence of at least 80% identity to residues 22-122 of the present sequence, or a sequence encoded by a nucleic acid hybridising under stringent conditions to the complement of the coding region comprising 1031 nucleotides, and having at least one biological activity of the polypeptide encoded by clone P00210009. The polypeptides and polynucleotides of the invention are useful for the treatment of activity creal and inflammatory diseases. The polynucleotides are useful in antisense mediated gene inhibition and in gene therapy. The polypeptides are useful in assays for identifying lead compounds that may be used as therapeutic agents in the treatment of cardiac, kidney or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat; secreted factor polypeptide; cardiac disease; renal disease; kidney; inflammatory disease; congestive heart failure; myocarditis; asthma; ss; dilated congestive cardiomyopathy; angina pectoris; cardiac arrhythmia; myocardial infarction; pulmonary hypertension; arteriosclerosis; stroke; atherosclerosis; cardiac tumour; glomerulonephritis; nephrotic syndrome; renal infarction; hereditary nephritis; polycystic kidney disease; chronic renal failure; renal vain thromboais; medullary sponge kidney; rheumatoid arthritis; ostcoarthritis; psoriasis; restenosis; PCR primer; graft versus host reaction; Crohn's disease; ulcerative colitis; probe; Alzhelmer's disease; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                            present sequence corresponds to polylinker DNA of the phagemid vector
                                                                                                                                                                                                                                                      Novel secreted factor encoded by clone P00210D09 useful for diagnosing, treating and/or preventing various cardiac, renal and inflammatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                factor DNA oligonucleotide probe #6.
                                                                                                                                                                                                                                                                                                                                      Example 1; Page 41; 69pp; English.
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                                                       27-SEP-2000; 2000WO-US026582.
                                                                                             99US-0156277P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS94743 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18; Conservative
                                                                                                                                                                           Stanton LW, Kapoun AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inflammatory diseases
                                                                                                                                                                                                              WPI; 2001-328177/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                      (SCIO-) SCIOS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200174901-A2
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                                                                                               27-SEP-1999;
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                05-APR-2001.
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                                                                                                                                                                                                                                                                                                 diseases
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AAS94743/c
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Gaps

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The invention relates to rat secreted factor polypeptides and the polynucleotides encoding them. The sequences are useful for treating cardiac, renal or inflammatory diseases. These include cardiac diseases such as congestive heart failure, myocarditis, dilated congestive cardiomyopathy, angina pectoris, myocardial infarction, cardiac cardiac tumours, renal diseases such as glomerulonephritis, nephrotic syndrome, renal infarction, hereditary nephritis, polycystic kidney disease, chronic renal failure, renal vein thrombosis and medullary consumptivitis, and infarmatory diseases such as asthma, rheumatoid arthritis, arthritis, stroke, psoriasis, restenosis, graft versus chost reaction, Crohn's disease, ulcerative colitis and Alzheimer's disease. Sequences AAS94693-AAS9475 represent cDNA clones, which encode the secreted factor polypeptides of the invention, and oligonucleotide
                                                                                                                                                                                                                                        Novel secreted factor polypeptide useful for treating cardiac diseases such as arteriosclerosis, myocardial infarction, inflammatory diseases such as asthma, stroke, and rheumatoid arthritis and renal diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 Example 1; Page 51; 189pp; English
23-MAR-2001; 2001WO-US009555.
                                           31-MAR-2000; 2000US-0193548P. 14-MAR-2001; 2001US-00809545.
                                                                                                                                                          Stanton LW, White RT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probes and PCR primers
                                                                                                                                                                                                 WPI; 2002-010779/01.
                                                                                                           (SCIO-) SCIOS INC.
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Gaps , 0 0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; ive 0; Mismatches 0; Indels Conservative Local Similarity 18; Query Match Matches

2709 AAAAAAAAAAAAAA 2726 18 AAAAAAAAAAAAAA 1 g ઠે

Angiogenesis inhibitory oligonucleotide #939. ABS78455 standard; DNA; 18 (first entry) 13-DEC-2002 ABS78455; RESULT 1063 

BP.

Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar

WO200253141-A2 Synthetic

11-JUL-2002.

14-DEC-2001; 2001WO-US048458.

14-DEC-2000; 2000US-0255534P.

(COLE-) COLEY PHARM GROUP INC.

Bratzler RL;

WPI; 2002-566690/60

Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.

Claim 2; Page 36; 276pp; English.

administering at least one antiangingenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangingenic nucled acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, obler-Webber Syndrome, myocardial angiogenesis, plaque meovascularisation, telangiectasia, hemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic The invention relates to inhibiting angiogenesis in a subject, acid of the invention

Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

Gaps ö 0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; ive 0; Mismatches 0; Indels 18; Conservative Local Similarity Query Match Matches

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ABS78429 standard; DNA; 18 (first entry) 13-DEC-2002 ABS78429; RESULT 1064 ABS78429/ 

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BP.

Angiogenesis inhibitory oligonucleotide #913.

Anglogenesis inhibitor; ss; anglogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophilac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.

Synthetic.

WO200253141-A2.

11-JUL-2002.

14-DEC-2001; 2001WO-US048458.

14-DEC-2000; 2000US-0255534P.

(COLE-) COLEY PHARM GROUP INC

Bratzler RL;

WPI; 2002-566690/60

Inhibiting angiogenesis in a subject, involves administering at least one

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The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject container acids acids, and instructions for administering them to a subject thaving a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque convoluarisation, intestinal adhesions, atheroselerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                                                                                                                                                                                                                                     0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
tive 0; Mismatches 0; Indels
antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid SEQ ID NO: 837.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ocation/Qualifiers
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/*trag= a
'~d base= OTHER
"whosphorc
                                 Page 35; 276pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                        ABL39401 standard; DNA; 18 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                         18; Conservative
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                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       developing cancer
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modified_base
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                                   Claim 2;
                                                                                                                                                                                                                                                                                                                                          Query Match
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The present invention relates to methods for treating or preventing

Disclosure; Page 308; 312pp; English.

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                                                                                                                                                                                                                                     the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful for destroying tumors, for identifying novel targets for the development of anti-tumor agents, and as specific ion channel modulators.
                               expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Apoptosis; ion channel modulator; hyperproliferative disease; tumour; therapy; leukaemia; carcinoma; sarcoma; degenerative disease; melanoma; Alzhaimer's disease; parkinson's diseases arteriosclerosis; melanoma; heart disease; stroke; vascular disease; nootropic; neuroprotective; cerebroprotective; cardiant; cytotoxic protein; cyplasin L; ss.
cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expressive cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cerophageal cancer, colon and rectum cancer, connective tissue cancer, nesophageal cancer, ingressive cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, spancreatic cancer, prostate cancer, habdomysaarcoma, skin cancer, stomach cancer, resticular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in texemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel polypeptide having cytotoxic activity obtainable from Aplysia, useful for destroying tumors, for identifying novel rarners for the
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                                                                                                                                                                                                                                                                                                                                       Length 18;
                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                               BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rudel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD41497 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                      18; Conservative
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                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1066
                                                                                                                                                                                                                                                                                                                                                                                      Matches
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preferably anti-tumour agents and as specific ion channel modulators, e.g., blockers or openers for therapy, diagnostic or research. They are useful for the diagnosis and therapy of hyperproliferative diseases, preferably tumours, e.g., leuksemia, carcinoma, sarcoma and melanoma. They are also useful for development of drugs for the treatment of degenerative diseases such as Alzheimer's disease, Parkinson's disease, arteriosclerosis, heart diseases, stroke and vascular diseases. The present sequence is an oligonucleotide which is used for amplifying sea hare cyplasin L DNA. This sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oĘ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a novel method for increasing the efficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Increasing efficiency of second strand cDNA synthesis using terminal continuation model before performing further RNA amplification by RNA
                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          continuation; TC; ss; second strand cDNA synthesis; primer;
                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                               Length 18;
                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                         Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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100.0%; Pred. No. 8.7e+02;
iive 0; Mismatches 0;
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RES FOUND MENTAL HYGIENE INC.
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2001US-0268664P.
2001US-0306216P.
2001US-0344557P.
2001US-0348242P.
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                                                                                                                                                                                                                                                                                                                                                                                                        437/c
ABS53437 standard; DNA; 18
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                                                                                                                                                                                                                                           Local Similarity 100.
1es 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Poly d(T) primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200265093-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-FEB-2001; 2
14-FEB-2001; 2
18-JUL-2001; 2
07-NOV-2001; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Terminal co
                                                                                                                                                           invention
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                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS53437;
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The present invention describes a method for the comparative detection of the amount of an RNA. The method comprises: (a) CDNAs obtained by transcribing respectively from at least two tissue RNAs are respectively from at least two tissue RNAs are respectively fragmented by using a same restriction enzyme; (b) each different adaptor and a common adaptor are added to each of the cDNA fragments derived from added cDNAs are mixed together; (d) and appror primer having the common sequence to said different adaptor and a gene specific adaptor are used to amplify said adaptor-added cDNAs containing no region derived from polyadenylic acid of the mRNA before the addition of the adaptor among the adaptor-added cDNAs prepared by the step (b); (e) the ratios of the cDNA amounts are measured between the tissues; (f) the RNA is detected from the measured result; (g) each different adaptor and a common adaptor are added to each of the genomic DNA fragments derived from a same or different individuals; (h) the resultant adaptor-added genomic DNAs are
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of the invention is useful for increasing the efficiency of second strand CODA synthesis and may be used for linear amplification of genetic signals from histologically stained tissue. The present sequence represents a poly d(T) PCR primer used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mixed together; (i) the adaptor-added genomic DNAs are amplified by using an adaptor primer having the common sequence to the different adaptor and a sequence-specific adaptor; and (j) the ratios of the amplified amounts of the genomic DNAs are measured between the individuals. The method is
                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     used for the detection of the amounts of RNA and DNA. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    represents an oligonucleotide which is used in the
                                                                                                                                          0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Comparative detection of the amounts of RNA and DNA.
                                                                                                     Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection; comparative detection; adaptor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Adaptor oligonucleotide SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 9; 9pp; Japanese.
                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                        ABA93239 standard; DNA; 18
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                                                                                                                                                                                       Conservative
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Matches 18; Conserv
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Best Local Similarity
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                                                                                                                                          Query Match
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Acyclic linker;
                   antisense; ss
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                                       Unidentified.
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                                                                                 08-MAY-2003
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                                                                                                                                                                    Damha MJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to an acyclic linker-containing oligonucleotide comprising at least one modified decoxyribonucleotide. Oligonucleotides of rehe invention are useful for preventing or decreasing translation.

They are useful for selectively preventing on a target RNA in a system. They are useful for selectively preventing gene expression in a sequencement for viral RNA, to hybridising to complementary RNA such as cellular mRNA or viral RNA, to hybridise to and induce cleavage of complementary RNA. They are also useful therapeutically in formulations or medicaments particular target RNA. The invention is used in gene therapy. The present sequence is a target RNA, used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                       Novel acyclic linker-containing oligonucleotide useful for preventing or decreasing translation, reverse transcription and/or replication of a target RNA in a system, comprises a modified deoxyribonucleotide.
Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense oligo #1, to elicit RNase H degradation of target RNA
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                                                                                                                                                           Target RNA #1 used in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                               Min K;
Indels
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                                                                                                                                                                                                                                                                                                                               Parniak MA,
                                                                                                                                                                                Acyclic linker; gene expression; gene therapy; ss
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                Mangos MM,
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                   2726
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Best Local Similarity 100.0%; Pr
Matches 18; Conservative 0;
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                                                                                             AAD56466 standard; RNA; 18 BP.
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                                        18 AAAAAAAAAAAAAAAA
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AAD56440 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                Damha MJ, Viazovkina E,
 18; Conservative
                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-421516/39.
                                                                                                                                                                                                                                                                                                           (UYMC-) UNIV MCGILL.
                                                                                                                                                                                                                         WO2003037909-A1.
                                                                                                                                                                                                     Unidentified.
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                                                                                                                                       07-AUG-2003
                                                                                                                                                                                                                                             08-MAY-2003
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                                                                                                                  AAD56466;
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                                                                         RESULT 1069
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Matches
  Matches
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The invention relates to an acyclic linker-containing oligonucleotide comprising at least one modified decoxyribonucleotide. Oligonucleotides of the invention are useful for preventing or decreasing translation, reverse transcription and/or replication of a target RNA in a system. They are useful for selectively preventing gene expression in a sequence-specific manner, for hybridising to complementary RNA such as cellular mRNA or viral RNA, to hybridise to and induce cleavage of complementary RNA. They are also useful therapeutically in formulations or medicaments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to prevent or treat a disease characterised by the expression of a particular target RNA. The invention is used in gene therapy. The present sequence is an antisense oligo used to elicit human RNase (ribonuclease) H degradation of target RNA. This sequence is used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel acyclic linker-containing oligonuclectide useful for preventing or decreasing translation, reverse transcription and/or replication of a target RNA in a system, comprises a modified deoxyribonuclectide.
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gene expression; gene therapy; ribonuclease; RNase H;
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100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mangos MM, Parniak MA, Min K;
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Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Viazovkina
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WPI; 2003-521815/49.

Parniak MA,

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The invention relates to an acyclic linker-containing oligonucleotide comprising at least one modified deoxyribonucleotide. Oligonucleotides of the invention are useful for preventing or decreasing translation, reverse transcription and/or replication of a target RNA in a system. They are useful for selectively preventing gene expression in a sequence-specific manner, for hybridising to complementary RNA such as cellular mRNA or viral RNA, to hybridise to and induce cleavage of complementary RNA. They are also useful therapeutically in formulations or medicaments to prevent or treat a disease characterised by the expression of a particular target RNA. The invention is used in gene therapy. The present sequence is an antisense oligo used to elicit human RNase (ribonuclasse) and degradation of target RNA. This sequence is used in the exemplification
                                                                                                                                                                                                                                        decreasing translation, reverse transcription and/or replication of a target RNA in a system, comprises a modified deoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunostimulatory, antiinflammatory, dermatological, antipsoriatic, antiulcer; gene therapy, vaccine, non-allergic inflammatory disease, psoriasis, eczema, allergic contact dermatitis, latex dermatitis, inflammatory bowel disease, ulcerative colitis, Crohn's disease; ss.
                                                                                                                                                                                                                      Novel acyclic linker-containing oligonucleotide useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
cive 0; Mismatches 0; Indels
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                                                                                                                                                     Mangos MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunostimulatory nucleic acid #882.
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                                                                                 29-OCT-2001; 2001US-0330719P
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                                                                                                                                                     Viazovkina
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                                                                                                                   (UYMC-) UNIV MCGILL
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Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2003050268-A1.
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               08-MAY-2003
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                                                                                                                                                     Damha MJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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New oligonucleotide, useful for preventing or treating a disease related to a target RNA in a system, e.g., AIDS or hepatitis B.
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                                                                                                                                                            The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic context dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sugar-modified nucleoside, acquired immune deficiency syndrome, AIDS; hepatitis B; gene therapy; virucide; anti-HIV; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                  Treating non-allergic inflammatory diseases, such as psoriasis, ecze allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. .v.
                                                                                                                             Disclosure; Page 33; 229pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       2726
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                                                                                                                                                                                                                                                                                                                                                                            0.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.7
Best Local Similarity 100.
Matches 18; Conservative
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Krieg AM, Berg DJ;

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                                                                                                                                                                                                                                                                                                                                                                                 Sugar-modified nucleoside, acquired immune deficiency syndrome, ALDS; hepatitis B; gene therapy; virucide; anti-HIV; antisense; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                          Antisense DNA-RNA hybrid #2 used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide, useful for preventing or treating a disease related
to a target RNA in a system, e.g., AIDS or hepatitis B.
target RNA in a system or preventing or treating a disease related to a target RNA in a system, e.g., acquired immune deficiency syndrome (AIDS) or hegatitis B. The invention is useful in gene therapy. The present sequence is an antisense oligonucleotide used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a new oligonucleoside which comprises alternating first and second segments. The first segment comprises at least one sugar modified nucleoside. The second segment comprises at least one 2'-deoxynucleoside. The oligonucleoside comprises at least 2 cach of the first and second segments, so that it comprises at least 4 alternating segments. The oligonucleotide is useful for preparing a composition for inducing RNase H-mediated cleavage of a target RNA in a
                                                                                                                                            Gaps
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                                                                                                                Length 18;
                                                                                                                                          0; Indels
                                                                                   Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                               DB 1; Le
8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "2'-0-methyl-D-uridine"
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                                                                                                             0.7%; Score 18; DB
100.0%; Pred. No. 8.7
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                       BP.
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/*tag= b
/label= RNA
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/label= RNA
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/*tag= a
/label= RNA
                                                                                                                                                                                                                                                                        AAD57878 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                            Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .15
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                    AAD57878;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sugar-modified nucleoside; acquired immune deficiency syndrome; AIDS; hepatitis B; gene therapy; virucide; anti-HIV; antisense; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense DNA-RNA hybrid #3 used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligonucleotide, useful for preventing or treating a disease related to a target RNA in a system, e.g., AIDS or hepatitis {\tt B}.
replication of a target RNA in a system, detecting the presence of a target RNA in a system, detecting the presence of a target RNA in a system, validating a gene target corresponding to a target RNA in a system, or preventing or treating a disease related to a target RNA in a system, e.g., acquired immune deficiency syndrome (AIDS) or hepatitis B. The invention is useful in gene therapy. The present
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                                                                                                                                                                                    Score 18; DB 1; Length 18; Pred. No. 8.7e+02;
                                                                                                                                                                                                                       0; Indels
                                                                                                                                                     Sequence 18 BP; 0 A; 0 C; 0 G; 9 T; 9 U; 0 Other;
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/*tag= b
//label= RNA
/note= "2'-0-methyl-D-uridine"
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/note= "2'-0-methyl-D-uridine"
                                                                                                                                                                                                    100.0%; Pred. No. 8., tive 0; Mismatches
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                                                                                                                                                                                                                                                                               BP.
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                                                                                                                                                                                                                           18; Conservative
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                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                             the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
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RESULT 1077
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                                                                                                                                                                                                                     Sugar-modified nucleoside, acquired immune deficiency syndrome, AIDS, hepatitis B; gene therapy; virucide; anti-HIV; antisense; DNA-RNA hybrid;
replication of a target RNA in a system, detecting the presence of a target RNA in a system, validating a gene target corresponding to a target RNA in a system or preventing or treating a disease related to a target RNA in a system, e.g., acquired immune deficiency syndrome (AIDS) or hepatitis B. The invention is useful in gene therapy. The present sequence is an antisense DNA-RNA hybrid used in the exemplification of
                                                                                                                                                                                                       Antisense DNA-RNA hybrid #1 used in the exemplification of the invention
                                                                                           Gaps
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                                                                         Sequence 18 BP; 0 A; 0 C; 0 G; 6 T; 12 U; 0 Other;
                                                                                                                                                                                                                                                                                 *tag= a
|abel= RNA
note= "2'-0-methyl-D-uridine"
                                                                                                                                                                                                                                                                                                             *tag= b
|abel= RNA
note= "2'-O-methyl-D-uridine"
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/label= RNA
/note= "2'-O-methyl-D-uridine"
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|abel= RNA
note= "2'-O-methyl-D-uridine"
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|abel= RNA
note= "2'-O-methyl-D-uridine"
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                                                                                                                 18 AAAAAAAAAAAAAA 1
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RNA
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label= RNA
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|abel= RNA
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RNA
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label= !
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label=
                                                                                 Best Local Similarity
Matches 18; Conserv
                                              the invention
                                                                                                                                                                                                                                                   Unidentified
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                                                                                                                                            RESULT 1076
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The present invention relates to a new oligonucleoside which comprises alternating first and second segments. The first segment comprises at least one sugar modified nucleoside. The second segment comprises at least of least one 2-deoxynucleoside. The oligonucleoside comprises at least 2 cach of the first and second segments, so that it comprises at least 4 alternating segments. The oligonucleotide is useful for preparing a composition for inducing RNAse H-mediated cleavage of a target RNA in a composition of a target RNA in a system, transcription or target RNA in a system, and spaced corresponding to a target RNA in a system or preventing or treating a disease related to a target RNA in a system, or preventing or treating a disease related to a target RNA in a system, or preventing or treating a disease related to a target RNA in a system, or useful inn gene therapy. The present corresponding to a target RNA in a system, or useful inn gene therapy. The present corresponding to a target RNA in a system, a system or useful in gene therapy. The present corresponding to a target RNA in a system or bepatities an antisense DNA-RNA hybrid used in the exemplification of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide, useful for preventing or treating a disease related to a target RNA in a system, e.g., AIDS or hepatitis B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 35; 73pp; English.
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                  Parniak MA;
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WO2003064441-A2.
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                                                                       07-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                  Damha MJ,
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                                                                                           The present invention relates to a new oligonucleoside which comprises alternating first and second segments. The first segment comprises at least one sugar modified nucleoside. The second segment comprises at least cach of the first and second segments, so that it comprises at least 2 of each of the first and second segments, so that it comprises at least 4 alternating segments. The oligonucleotide is useful for preparing a composition for inducing RNase H-mediated cleavage of a target RNA in a system, preventing or decreasing translation, transcription or replication of a target RNA in a system, detecting the presence of a target RNA in a system, of the target RNA in a system, and the segment of a target RNA in a system, of treating a disease related to a target RNA in a system, of treating a disease related to a target RNA in a system, e.g., acquired immune deficiency syndrome (AIDS) or hepatitis B. The invention is useful in gene therapy. The present a sequence is a target RNA used in RNase H assay. This sequence is used in
                                                                                                                                                                                                                                                                                                                                                                           ö
                            New oligonucleotide, useful for preventing or treating a disease related to a target RNA in a system, e.g., AIDS or hepatitis B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fouron Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory nucleic acid #824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 17; 221pp; English.
                                                                                                                                                                                                                                                                                              the exemplification of the invention
                                                                       Example 4; Page 38; 73pp; English
                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB37210 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-DEC-2003 (first entry)
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BRAT/) BRATZLER R L.
(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-657977/62.
  WPI; 2003-689523/65.
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invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ds; allergy; asthma; poly-G nucleic acid; aerosol formulation; hypo-responsive subject; immunostimulatory.
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8; Conservative 0; Mismatches 0; Indels
                                                                                                                                               Query Match
0.7%; Score 18; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 8.7e+02;
Matches 18; Conservative 0; Mismatches 0; Indels
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                                                                                                   Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 18; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulatory nucleic acid #850.
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                                                                                                                                                                                                                                                                                      ADE77617 standard; DNA; 18
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(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-657977/62
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ADB37236/c
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Matches
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ADE77617;

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The present invention relates to a method for detecting nucleic acid (NA). The method involves hybridizing the NA with nucleic acid target sequence obtained from reverse transcription or RNA amplification using the sample RNA as a template and single stranded probes immobilized on a support body. The present sequence was used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detecting nucleic acid, by hybridizing nucleic acid with other nucleic acid target obtained from reverse transcription or RNA amplification, an other single stranded nucleic acid probe immobilized on a support body.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid amplification; RNA transcription; RNA polymerase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                           DNA detection; RNA detection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleotide sequence of an oligo dT18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 6; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-2001; 2001JP-00309382.
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                                                                   (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                (HITA ) HITACHI LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the invention.
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                                                                   16-JUN-2005
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                                                                                                                 Primer #10
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                         ADZ47933;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for the concurrent interrogation of a number of polymorphic sites in the presence of, and without interference from, non-designated polymorphic sites. Specifically, it comprises conducting a multiplexed elongation assay by applying one or more temperature cycles to achieve linear amplification of the target or a combination of annealing and elongation steps under temperature. Controlled conditions. Furthermore, this detection method uses probe extension or elongation and relies on enzymatic recognition, a superior cechnique that no longar depends on differential hybridisation. The present invention describes probes and methods useful for identifying or detecting polymorphisms at one or more designated sites, such that they can identify mutations within the cystic fibrosis conductance transmembrane regulator (GFTR) or the human leukocyte antigen (HLA) genes. In addition, concurrent interrogation of a multiplicity of genetic profiling, and identity testing. This classified is the negative control probe used for the elongation mediated multiplexed analysis of HLA-DR, in an exemplification of the
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                                                                                                                             probe; ss; negative control; CFTR; human leukocyte antigen; HLA; genetic testing; carrier screening; genotyping; profiling; polymorphic; multiplexed elongation assay; enzymatic recognition; cystic fibrosis conductance transmembrane regulator.
                                                                                      Human probe NEG for elongation mediated multiplexed analysis of HLA-DR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Concurrent interrogation of a number of polymorphic sites, useful for genetic testing, carrier screening, genetic profiling, and identity testing, comprises conducting a multiplexed elongation assay using
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15-OCT-2001; 2001US-0329629P.
15-OCT-2001; 2001US-0329620P.
14-MAR-2002; 2002US-036441EP.
                                                                                                                                                                                                                                                                                                                                                                                                      15-OCT-2002; 2002WO-US033012.
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                                       29-JAN-2004 (first entry)
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                          Synthetic.
Homo sapiens.
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Gaps

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Kamme FC, Zhu JY;

7933/c ADZ47933 standard; DNA; 18 BP.

RESULT 1081 ADZ47933/c ID ADZ47

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invention.

WPI; 2004-035466/03.

fluid in a chamber comprising introducing into the chamber a support to

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                                                                                                                     The invention relates to amplifying for RNA in a sample comprises eliminating single-stranded oligonucleotide from the transcription sample. The method involves synthesizing single-stranded cDNA by incubating the sample RNA with reverse transcriptase and an oligonucleotide primer that primes synthesis in a direction toward 5' end of the RNA; converting the single-stranded cDNA into double-stranded cDNA to form a transcription sample containing a cDNA template; eliminating single-stranded oligonucleotide from the transcription sample; and transcribing the cDNA template into RNA using an RNA polymerase. The method is useful for improving RNA polymerase based RNA transcription from a polynucleotide template. The method inhibits the undesired nontemplate derived production of RNA in the transcription reaction. Sequences ADI34483-ADI34489 represent oligonucleotides used in a T7 RNA
                          Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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fluid reagent distribution; time dependent fluid reactivity; ss.
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100.0%; Pred. No. 8.7e+02;
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                                                                                            Example 1; SEQ ID NO 8; 26pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rest element oligonucleotide #2.
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                     ranscription reaction.
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(AMOR/) AMORESE D A.
(PECK/) PECK B J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JS2003232343-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH78590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1083
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The invention relates to a method of determining a functional property of

Example 2; SEQ ID NO 2; 22pp; English.

Determining a functional property of fluid in chamber by introducing a support comprising test elements having reaction and detection domains, introducing a test fluid, and detecting locations not reactive with the

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Peck

Amorese DA,

Leproust EM,

WPI; 2004-061269/06.

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which is bound several test elements, each of the test elements comprising a reaction domain and a detection domain, introducing into the chamber a fluid that is interactive with the reaction domains, removing the fluid from the chamber, determining by means of the detection domains, removing the locations at which the fluid has not interacted with the reaction domains the locations to the functional property of the comprises a member of a specific binding pair. The detection domain comprises a member of a specific binding pair. The detection domain comprises a member of a specific binding pair. The determining of the comprises interacted with the fluid. The functional property is chosen from the flow pattern of the fluid, respect the fluid and time dependent reactivity of the fluid. The method is useful for determining a functional property of a fluid in a chamber and for synthesising arrays of biopolymers e.g., arrays of polymucleotides. The method provides for the characterisation of a new fluid in a known flow call, a known fluid in a new flow call or a new fluid in the method combination. This sequence represents a test element used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a method of synthesising single stranded cDNA by a 5'-ligated adaptor-mediated process involving: annealing a CDNA synthesis primer to RNA, separating the cDNA strand from the RNA, purifying the cDNA, contacting the cDNA with an adaptor, ligating the adaptor through 5'-phosphate on strand (II) of the adaptor to single stranded using DNA ligase, and amplifying the obtained ligated single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthesizing single stranded cDNA, involves annealing cDNA synthesis primer to RNA and synthesizing first cDNA strand, ligating adaptor to single stranded cDNA using DNA ligase, and amplifying ligated single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single stranded cDNA; adaptor-mediated process; cDNA synthesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single stranded cDNA production poly-A-tail seqid 6.
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pettersson NB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 6; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2709 AAAAAAAAAAAAAAA 2726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-AUG-2000; 2000US-0226954P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (AZIG-) AZIGN BIOSCIENCE AS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO28710 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 1000.
Matches 18, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             stranded cDNA fragment.
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                                                                                                                                                                                                                                                                                                                                                                     of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            poly-A-tail; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rhirstrup K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6706476-B1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADO28710;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1084
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$$$$$$$$$$$$$$$$$$$$$$
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cc stranded fragment in an molecular amplification procedure. The method is useful for: synthesising a single stranded cDNA by a 5'-ligated adaptor-cc mediated process, where the source of nucleic acid is chosen from blood, serum, plasma, cerebrospinal fluid, urine, tissue samples, biopsies and caliva. The tissue sample comprises a cell population which may be single cell, 100-1000000 cells or more as desired, making a cDNA library from a collection of mRNA molecules in a sample, where the method is applied to amplify the cDNAs corresponding to the mRNAs by annealing one or more convariant of mRNA molecules in a sample, where the method is applied to amplify the cDNAs corresponding to the mRNAs by annealing one or more convariant is modified by by biotin in the 5' end, isolating the biotin-containing single stranded cDNA (sense) by use of streptavidin coated containing single stranded cDNA (sense) by use of streptavidin coated containing single stranded cDNA (sense) by use of streptavidin coated magnetic beads, hybridising the sense to the anti-sense cDNA, where an unhybridised sub-population of the anti-sense cDNA is found, isolating the unhybridised sub-population of the anti-sense cDNA, where an unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population by PCR using primer I and correction from the unhybridised sub-population by PCR using primer I and collection from the unhybridised sub-population of the anti-sense cDNA is primer. The method is an improved method for producing amplified cold class and containing application. The method is an improved method for producing amplified cold class and contains a poly-A-tail used to in the contains of cDNA colds.
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Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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Gaps
                                          ó,
               Length 18;
                                          0; Indels
               Score 18; DB 1; Le
Pred. No. 8.7e+02;
Query Match 0.7%; Score 18; ub .
Best Local Similarity 100.0%; Pred. No. 8.7.
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ઠે g ADO28711 standard; DNA; 18 BP AD028711;

Single stranded cDNA production poly-A-tail complement segid 7. single stranded cDNA; adaptor-mediated process; cDNA synthesis; 15-JUL-2004 (first entry) poly-A-tail; ss. 

Synthetic.

16-MAR-2004.

US6706476-B1.

09-MAR-2001; 2001US-00803263.

22-AUG-2000; 2000US-0226954P.

(AZIG-) AZIGN BIOSCIENCE AS.

Thirstrup K, Warthoe P, Pettersson NB; WPI; 2004-326403/30. Synthesizing single stranded cDNA, involves annealing cDNA synthesis primer to RNA and synthesizing first cDNA strand, ligating adaptor to single stranded cDNA using DNA ligase, and amplifying ligated single stranded cDNA fragment.

Example 1; SEQ ID NO 7; 22pp; English.

The Invention describes a menthod of synthesising single stranded cubn by contracting describing the cDNA strand from the RNA, synthesis primer to RNA, separating the CDNA strand from the RNA, puritying the cDNA, sontacting the CDNA with an adaptor, ligating the adaptor through 5'-phosphate on strand (II) of the adaptor to single stranded using DNA ligase, and amplifying the obtained ligated single correspond a single stranded frasedure. The method is consent for synthesising a single stranded cDNA by a 5'-ligated adaptor-mediated process, where the source of mucleic acid is chosen from blood, serum, plasma, cerebrospinal fluid, urine, tissue samples, bioppies and saliva. The tissue sample comprises a cell population which may be single coll; 100-1000000 cells or more as desired; making a cDNA library from a collection of mRNA molecules in a sample, where the method is a spile stranded colls or more several mRNAs by annealing one or more constity the CDNAs corresponding to the mRNAs by annealing one or more constaining single stranded cDNA senses synthesisisng a double-stranded cDNA collection from a first mRNA population by the method, where primer 1 is modified by biotin in the 5' end, isolating the biotin-containing single stranded cDNA (sense) by use of streptavidin coated magnetic beads, synthesising a double-stranded cDNA, collection from a second mRNA population of the anti-sense cDNA, where an unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population of the anti-sense cDNA, collection from the unhybridised sub-population and preselected c invention describes a method of synthesising single stranded cDNA by used to in the production single stranded cDNA. X8666666666666666666666666666666666688X

Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

.. 0 0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; 0; Indels 100.0%; Pred. No. Query Match 0.7° Best Local Similarity 100. Matches 18; Conservative

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Gaps

2709 AAAAAAAAAAAAAA 2726 g 8

Synthetic leader sequence encoding DNA SEQ ID NO:77. ADO26684 standard; DNA; 18 BP. (first entry) 12-AUG-2004 AD026684; RESULT 1086 AD026684/c 

phenotype; phenotypic preference; phenotype modulation; leader; ds. Synthetic.

WO2004042059-A1. 21-MAY-2004. 10-NOV-2003; 2003WO-AU001487. 08-NOV-2002; 2002US-0425163P.

(UYQU ) UNIV QUEENSLAND.

Frazer IH;

WPI; 2004-411519/38. P-PSDB; ADO26685.

21-MAY-2004.

The present invention describes a method for constructing a synthetic polyuncleotide from which a polypepide is producible to confer a selected phenotype to an organism of interest or part in a different colorate polymorlaceide for minitoh a polypepide is producible to confer a selected phenotype to an organisms of interest a different polymorlaceide for replacement with a synonymous codon of the parent polymorlaceide for replacement with a synonymous codon of the parent polymorlaceide for replacement with a synonymous codon of the selected from organisms of parts, where the test organism are selected from organisms of the same species as the organism of interest and organisms that are related to the organisms of interest confined to phenotypic preference of a first codon with the synonymous codon to construct the synthetic polymorlaceided. Also described: (1) a method for che synthetic polymorlaceided. Also described: (1) a method for interest or its parts; (2) a synthetic polymorlaceide constructed from the method above, (3) an organism or interest or part containing a synthetic polymorlaceide constructed from the method above, (4) an organism or interest or part containing a synthetic onstruct that comparises a resporter protein expension of a first dodon fued in frame with a reporter polymorlaceide operator of a selected phenotype that is displayed by an organism of interest or part and that results from the expression of a parent polymorlaceide that condess the polypeptide, and (7) a method of reducing the quality of a selected phenotype that is displayed by an organism of reducing that results from the ex Constructing synthetic polynucleotide for modulating the quality of a selected phenotype displayed by an organism comprises replacing a first codon with a synonymous codon to construct the synthetic polynucleotide. Example 1; SEQ ID NO 77; 86pp; English.

Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; 0; Indels 100.0%; Prec. ... 2709 AAAAAAAAAAAAAAAA 2726 18; Conservative Query Match Best Local Similarity Matches ઠ 셤

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18 ААААААААААААААА 1

ADO26682 standard; DNA; 18 BP 12-AUG-2004 (first entry) ADO26682; RESULT 1087 ADO2668; 

Synthetic leader sequence encoding DNA SEQ ID NO:75.

phenotype; phenotypic preference; phenotype modulation; leader; ds.

Synthetic.

WO2004042059-A1.

The present invention describes a method for constructing a synthetic polymucleotide from which a polypeptide is producible to confer a selected phenotype to an organism of interest or part in a different gality than that conferred by a parent polymucleotide from selected or conferred by a parent polymucleotide. The method comprises: (a) selecting a first codon of the parent polymucleotide for replacement with a synonymous codon of the parent polymucleotide for replacement with a synonymous codon of different phenotypic preference than the first codon in a comparison of phenotypic preference than the first codon in a comparison of construct phenotypic preference than the first codon in a comparison of interest and organisms that are related to the organisms of interest of interest and organisms that are related to the organisms of interest of interest and organisms that are related to the organisms of interest or its parts: (2) a synthetic polymucleotide constructed from interest or its parts: (2) a synthetic polymucleotide constructed from interest or its parts: (2) a synthetic polymucleotide for organism or interest or part containing a synthetic construct that comparises a regulatory polymucleotide operator or part containing a synthetic construct that comparises a regulatory polymucleotide operator polymucleotide that conforms a predicted to produce a selected phenotype that is displayed by an organism of interest or part of a selected phenotype that is displayed by an organism of interest or part or part and that results from the expression of a parent polymucleotide that encodes the polypeptide; (6) a method of reducing the quality of a selected phenotype that is displayed by an organism of interest or part or encodes the polypeptide; and (7) a method of reducing the quality of a selected phenotype to an organism of a parent polymucleotide that encodes the polypeptide; and (7) a method of reducing the quality of a selected phenotype to an organism of interest or part in a displayed by an organism of salec Constructing synthetic polynucleotide for modulating the quality of a selected phenotype displayed by an organism comprises replacing a first codon with a synonymous codon to construct the synthetic polynucleotide. Example 1; SEQ ID NO 75; 86pp; English. 10-NOV-2003; 2003WO-AU001487. 08-NOV-2002; 2002US-0425163P. (UYOU ) UNIV QUEENSLAND WPI; 2004-411519/38. P-PSDB; ADO26683 Frazer IH; 

Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

ö 0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; tive 0; Mismatches 0; Indels Best\_Local Similarity 100. Matches 18; Conservative Query Match

2709 AAAAAAAAAAAAAAA 2726

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18

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Gaps

RESULT 1088 ADP86130/c (first entry)

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New nucleic acid molecules encoding rat KDR protein, useful for identifying inhibitors of KDR activity for treating cancer, ischemic ocular diseases, and inflammation.
                                                                                                                                           Rat; receptor tyrosine kinase; KDR; therapy; cancer; ischaemic ocular disease; proliferative retinopathy; inflammation; reverse transcription; RT; PCR; primer; ss.
                                                                                                       Rat KDR cytosolic domain cloning RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Page 30; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                         23-JAN-2004; 2004WO-US001928
                                                                                                                                                                                                                                                                                                                                                                                             29-JAN-2003; 2003US-0443335P
                                                                                                                                                                                                                                                                                                                                                                                                                                      (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-604429/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pan B,
                                                                                                                                                                                                                                  Rattus norvegicus
                                                                                                                                                                                                                                                                         WO2004070004-A2.
                                                                   04-NOV-2004
                                                                                                                                                                                                                                                                                                               19-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thomas RA,
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ADR57967/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a class of CpG immunostimulatory
oligonuclectides containing a 5'TCG motif or a CG at or the 5' end that
are useful for stimulating an immune response. Oligonuclectides and
compositions of the invention are useful for treating allergy or asthma,
viral and bacterial infections and cancer e.g. biliary tract cancer,
or breast cancer, cervical cancer, choriocarcinoma, colon cancer,
commonstrial cancer, cenner, lymphomas, intraepithelial neoplasms,
liver cancer, lung cancer (e.g. small cell and non-small cell), melanoma,
neuroblastomas, ovarian cancer, pancreatic cancer, prostate cancer,
cectal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer,
and CNS cancer, connective tissue cancer, oseophageal cancer, eye cancer,
chodgkin's lymphoma, larynx cancer, oral cavity cancer, skin cancer,
testicular cancer, as well as other carcinomas and sarcomas. The
invention is also useful in gene therapy. The present sequence is a CpG
invention is also useful.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                            CpG immunostimulatory oligonucleotide; immune response; allergy; asthma; viral infection; bacterial infection; cancer; lymphoma; intraepithelial neoplasm; melanoma; neuroblastoma; Hodgkin's lymphoma; carcinoma; sarcoma; gene therapy; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast cancer, cervical cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Uhlmann E;
                                                                                                                      CpG immunostimulatory oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; SEQ ID NO 1; 104pp; English.
                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunostimulatory oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Krieg AM, Jurk M, Vollmer J,
  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHARM GROUP INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-DEC-2002; 2002US-0432409P.
25-SEP-2003; 2003US-0506108P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-DEC-2003; 2003WO-US039775.
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ADP86130 standard; DNA; 18
                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (COLE-) COLEY PHARM GROUP (COLE-) COLEY PHARM GMBH.
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Best Local Similarity 100.
Matches 18; Conservative
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/*tag=
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                                                                                                                                                                                                                                                                                                        Key
modified_base
                                                                                09-SEP-2004
                                                                                                                                                                                                                                                                  Unidentified
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                                        ADP86130;
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Mcgaughey

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The invention relates to rat receptor tyrosine kinase (KDR) and its corresponding nucleic acid sequence. The nucleic acid molecules of the invention are useful for identifying compounds that modulate wild-type rat KDR activity to evaluate the safety and efficacy of specific inhibitors of KDR in rats. KDR inhibitors are useful for treating cancer, inchammation. The present sequence is a reverse transcription (RT) PCR primer used for cloning rat KDR cytosolic domain. This sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; signal amplification method; gene expression; reverse transcription; self-assembly reaction; DNA chip.
                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                            Score 18; DB 1; Length 18;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                       Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide #4 for signal amplification method.
                                                                                                                                                                                                                                                            100.0%; Pred. No. b.,
                                                                                                                                                                                                                                                                                                                          2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                                                                                                                                                                                            0.7%;
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ADR57967 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                     18; Conservative
                                                                                                                                                                                                                                                                   Local Similarity
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2709 AAAAAAAAAAAAAAA 2726

AAAAAAAAAAAAAA

18

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ADR32355/c ID ADR32355 standard; DNA; 18 BP.

RESULT 1089

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RESULT 1092
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                                                                                                                                                                                  expressed gene using reverse transcription reaction and a self-assembly reaction of forming a self assembly of oligonucleotide probes, thus improving detection sensitivity of the expressed gene in a DNA chip. (MI) is useful for signal amplification method (MI) for detecting expressed gene (claimed). (MI) improves detection sensitivity of the expressed gene in a DNA chip (claimed). (MI) does not require use of expressive enzymes and enables detection corresponding to the original RNA length or expression amount because of using neither linear amplification nor PCR. This sequence corresponds to a nucleotide used in the method of the
                                                                                                                                                                          invention relates to a signal amplification method (M1) for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Preparing amplified RNA (aRNA) sequences present in single stranded or made single stranded target polynucleotide(8), useful for detecting
                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplified RNA preparation; nervous system disorder; neurodegenerative disease; Parkinson's disease; Alzheimer's disease; multiple sclerosis; psychiatric disorder; schizophrenia; affective disorder; manic depression; lack of appetite control; attention deficit disorder; cancer cell detection; ss.
                                                                                                            using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amplified RNA (aRNA) preparation method-related RNA sequence #1
                                                                                                                                                                                                                                                                                                                                                        .;
0
                                                                                                           Signal amplification method for detecting expressed gene, by reverse transcription reaction and self-assembly reaction of
                                                                                                                                                                                                                                                                                                                                  0.7%; Score 18; DB 1; Length 18;
00.0%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                            illarity 100.0%; Pred. No. 8.7
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Enright
                                                                                                                                                      Disclosure; SEQ ID NO 4; 27pp; Japanese.
                                           (PALM-) PALMA BEEZ RES INST CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ma X,
                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAAAA 2726
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13-FEB-2004; 2004WO-JP001588
                     14-FEB-2003; 2003JP-00037212
                                                                                                                                                                                                                                                                                                                                                                                                                                                  .010/c
ADT55010 standard; RNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Salunga RC,
                                                                                                                                oligonucleotide probes.
                                                                 Fujikawa T;
                                                                                      WPI; 2004-642306/62
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Erlander MG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-OCT-2004
                                                                                                                                                                                                                                                                                           invention.
                                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADT55010;
                                                                 Usui M,
                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1091
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The invention comprises a method of preparing amplified RNA (aRNA)

c sequences present in one or more target polynucleotide that is single
stranded or made single stranded. The method involves forming double
c stranded or made single stranded. The method involves forming double
c stranded cDNA templates containing sequences present in the target
c polynucleotide and transcribtion via the promoter region to produce
c amplified RNA containing sequences of the target polynucleotide. The
c method of the invention is useful for amplifying the population of RNAs
c spice variants. The method is also useful for determining gene
c extracted from formalin-fixed tissues and/or the population of RNAA
c splice variants. The method is also useful for determining gene
c extracted from in neuronal and non-neuronal cells involved in disorders of
the nervous system, such as: neurodegenerative diseases (e.g. Parkinson's
c disease, Alzheimer's disease, and multiple sclerosis); psychiatric
c disease, Alzheimer's disease, and multiple sclerosis); psychiatric
c depression, lack of appetite control, and attention deficit disorder).
C depression, lack of appetite control, and attention deficit disorder).
C fromethod of the invention may also be used to detect cancer cells, and
c of acilitate diagnosis/progrosis of cancer in a patient. The present RNA
c sequence is shown in a figure exemplifying the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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comprises transcribing double stranded cDNA templates with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amplified RNA (aRNA) preparation method-related DNA sequence #4
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 0 A; 0 C; 0 G; 0 T; 18 U; 0 Other;
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                                                                                                                            Disclosure; Fig 2; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 AAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADT55008 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-710328/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             an RNA polymerase
                                                        polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2004085681-A2
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The present invention relates to compositions for inhibiting influenza infection and/or replication. The compositions comprise a short interfering RNA (SiRNA) targeted to a target transcript, which is an agent specific transcript involved in infection by or replication of influenza A or B virus. The target transcript
                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 107; 241pp; English.
                                                                                                                                                                                                                                                                                                                                                                       Eisen HN,
                                                                                                                                                                                                                                                                                                  WO2004028471-A2.
                                                                                                                                                                                                                                                                                                              08-APR-2004.
                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                        ADN30833;
                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                       Chen J,
                                                                                                                                                   Matches
X3555555555555555555555
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The invention comprises a method of preparing amplified RNA (aRNA)

Sequences present in one or more target polymuclectide that is single
stranded crm made single stranded. The method involves forming double
stranded cDNA templates containing sequences present in the target

CC polymuclectide and transcribtion to the propertion of produce
cC amplified RNA containing sequences of the target polymuclectide. The

CC amplified RNA containing sugernic so it the target polymuclectide. The

CC amplified RNA containing sugernic so it the population of RNAs

CC capable of the invention is useful for amplifying the population of RNAs

CC extracted from formalin-fixed tissues and/or the population of RNAs

CC capable of the invention is useful for determining gene

CC expression in neuronal and non-neuronal cells involved in disorders of

CC the nervous system, such as: neurodegenerative diseases (e.g. Parkinson's

CC disease, Alzheimer's disease, and multiple sclerosis); psychiatric

CC disorders (e.g. schizophrenia); and affective disorders (e.g. manic

CC depression, lack of appetite control, and attention deficit disorder).

CC The method of the invention may also be used to detect cancer cells, and

CC decilitate diagnosis/prognosis of cancer in a patient. The present DNA

CC sequence is shown in a figure exemplifying the method of the invention. ö New composition comprising an siRNA or shRNA targeted to an agent-specific transcript, useful for treating or preventing influenza virus replication, pathogenicity or infectivity. Gaps ; 0 interfering RNA; siRNA; Score 18; DB 1; Length 18; Pred. No. 8.7e+02; 0; Indels Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other; Virucide; Vaccine; influenza virus infection; Mismatches influenza virus replication; short interfe short hairpin RNA; shRNA; ss; PCR; primer. (MASI ) MASSACHUSETTS INST TECHNOLOGY. 100.0%; Pre 2709 AAAAAAAAAAAAAAA 2726 1 AAAAAAAAAAAAAA 18 833/c ADN30833 standard; DNA; 18 BP. 28-SEP-2002; 2002US-0414457P. 10-FEB-2003; 2003US-0446377P. 29-SEP-2003; 2003WO-US030502 13-JAN-2005 (first entry) ö 18; Conservative ge PCR primer, SEQ 112. WPI; 2004-305101/28. Local Similarity

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The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-ga', and where the second dose is administered from 1 day to 8 weeks after the first dose. The methods and compositions of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, uricaria and atopic dermatitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suppressing allergies, including aethma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dos of an immunostimulatory nucleic acid.
              protein 1, membrane protein 2, nonstructural protein 1, nonstructural protein 2, polymerase protein PB1, polymerase protein PB2, polymerase protein PA or polymerase protein NP. The composition is useful for treating or preventing influenza virus replication, pathogenicity or infectivity. The present sequence is a PCR primer used during siRNA
                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; antiasthmatic; antiallergic; dermatological; antiinflammatory;
                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.
 encodes a protein consisting of hemagglutinin, neuraminidase,
                                                                                                                                                                          Score 18; DB 1; Length 18;
Pred. No. 8.7e+02;
                                                                                                                                                                                                            0; Indels
                                                                                                                                       Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Allergic response suppressor oligonucleotide #939.
                                                                                                                                                           0.7%; Scor.
100.0%; Pred. No. v...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-FEB-2000; 2000US-0179991P.
02-FEB-2001; 2001US-00776479.
                                                                                                                                                                                                                                                                                                                                                                      ADU90255 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Petersen DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                          18; Conservative
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                                                                                                                                                                                            Similarity
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(PETE/) PETERSEN D
(FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004235774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bratzler RL,
                                                                                                      preparation.
                                                                                                                                                                                                                                                                                                                                                                                                                                          10-FEB-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                              Local
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                                                                                                                                                                                                            Matches
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Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

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ADXS6134/c
ID ADXS6134 standard; DNA; 18 BP.
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                                                                                                                                                        Unidentified.
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24-FEB-2005
                                                                      21-APR-2005
                                                                                                                                                                                                         16-JUN-2004.
                                                                                                                                 primer; 88.
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                                            ADX56134;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of suppressing a symptom of an allergic response in a subject by administering a first and second dose of an immunostimulatory nucleic acid that comprises a nucleotide sequence comprising 5'-cg-3', and where the second dose is administered from 1 day present invention are useful for the treatment or prevention of the present invention are useful for the treatment or prevention of asthma and allergy, including rhinitis, urticaria and atopic dematitis, using an immunostimulatory nucleic acid alone or in combination with other medicaments. They can also be used in preventing bacterial and viral infections. This sequence represents an oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suppressing allergies, including asthma, rhinitis, urticaria and atopic dermatitis, in a subject, comprises administering a first and second dose of an immunostimulatory nucleic acid.
                                  Gaps
                                                                                                                                                                                                                                           se, antiasthmatic; antiallergic; dermatological; antiinflammatory; antibacterial; virucide; immunoglobulin E antagonist; allergy; immunostimulator; asthma; rhinitis; urticaria; dermatitis; bacterial infection; viral infection.
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           DB 1; Length 18;
                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                     Allergic response suppressor oligonucleotide #913.
                     8.7e+02;
hes 0;
                 100.0%; Pred. ...
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           0.7%; Score 18;
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                                                          2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                ADU90229 standard; DNA; 18 BP
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02-FEB-2001; 2001US-00776479.
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                                                                                     AAAAAAAAAAAAAAAAA
                                                                                                                                                                                                (first entry)
                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                              (BRAT/) BRATZLER R L.
(PETE/) PETERSEN D M.
(FOUR/) FOURON Y.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-833006/82.
                                                                                                                                                                                                                                                                                                                                   JS2004235774-A1.
                                                                                                                                                                                                10-FEB-2005
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                                                                                                                        RESULT 1095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vector and a recombinant vaccine composition against poliovirus, which may induce the formation of neutralizing antibodies against Sabin types 1, 2 and 3 polioviruses, and prevent the side-effects of OPV (attendated oral polio vaccine, Sabin). The present sequence is that of a PCR primer which was used during the development of the novel recombinant Sabin type 1 poliovirus vector of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Poly (dT)12-18 oligo, SEQ ID NO:150, used in first strand cDNA synthesis.
Novel recombinant Sabin type 1 poliovirus vector-related PCR primer #13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel recombinant Sabin type 1 poliovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                          vaccine; Sabin type 1 poliovirus; vector; poliovirus infection; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Recombinant sabin type 1 poliovirus vector and recombinant vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0; Indels
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/note= "Optionally and serially deleted"
                                                                                                                                                                                                                                                                                                                                                        Lee SG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                        Lee DS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense therapy; thioredoxin inhibitor; ss
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13. .18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                                                                               composition against poliovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 14; 32pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18 AAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                  10-DEC-2002; 2002KR-00078158
                                                                                                                                                                                                                                                                           10-DEC-2002; 2002KR-00078158.
                                                                                                                                                                                                                                                                                                                                                        Kim DY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADV11817 standard; DNA; 18
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Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                    (CREA-) CREAGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-715997/70.
                                                                                                                                                                                                                                                                                                                                                          Jung HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2004241717-A1
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18 AAAAAAAAAAAAAAA 1

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RESULT 1096

Tue Nov

ADW10182;

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10-FEB-2004; 2004US-00776933.
         10-FEB-2003; 2003US-0446374P
            (SANT-) SANTARIS PHARMA AS.
                Hansen B, Thrue CA,
 02-DEC-2004.
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Wissenbach M; Petersen KD, Westergaard M,

WPI; 2005-056301/06.

Novel compound useful for modulating expression of gene involved in cancer disease, or for modulating red blood cell proliferation, cellular proliferation, ion metabolism or glucose and energy metabolism.

Example 6; SEQ ID NO 150; 92pp; English.

peruchoises Selected in the 3-metropyraphracil, 6-aminopurine, 2aminopurine, inosine, diaminopurine and 2-chloro-6-aminopurine. The
invention also relates to a conjugate comprising a TRX antisense
coligonuclectide of the invention and one or more non-nuclectide or nonplarmaceutical composition comprising a TRX antisense oligonuclectide, and a
plarmaceutical composition comprising a TRX antisense oligonuclectide or
conjugate of the invention, optionally further comprising a
chemotherapeutic agent, an antiinflammatory compound or an antiviral
compound. The TRX antisense oligonuclectides, and conjugates and
compositions containing them, are useful in the treatment of cancers such
as carcinoma, breast carcinoma, are useful in the treatment of cancers such
as carcinoma, breast carcinoma, recurrent superficial bladder cancer,
carcinoma, bladder carcinoma, recurrent superficial bladder cancer,
stomach carcinoma, prostatic carcinoma, pancreatic carcinoma, lung
carcinoma, cervical carcinoma, colorectal carcinoma and carcinoid
tumors); sarcomas (e.g., osteosarcoma, Eshigs sarcoma, chondrosarcoma,
malignant fibrous histiccytoma, fibrosarcoma, and Kaposi's sarcoma;
creationant of the TRX antisense oligonuclectides are also useful in the
creationant of conditions such as atheroselerosis psoriasis, diabetic
creationant of conditions such as atheroselerosis psoriasis, diabetic
creationant of conditions such as atheroselerosis psoriasis, diabetic
creationant of conditions and as thereseleroselerome, and allarents. The invention relates to antisense oligonucleotides consisting of 8-50 nucleotides and/or nucleotide analogs which inhibit expression of the putative human oncogene thioredoxin (TRX). The antisense oligonucleotides comprise a subsequence of 8 or more nucleotides or nucleotide analogs, wherein the subsequence is located within a sequence selected from ADV11669-ADV11724. The oligonucleotides preferably contain at least nucleotide analog, especially a locked nucleic acid (INA) or a modified nucleobase selected from 5-methylcytosine, isocytosine, retinopathy, theumatoid arthritis, asthma, warts, and allergic dermatitis. They may additionally be used for inhibiting cellular proliferation and for modulating red blood cell proliferation, ion metabolism, glucose and energy metabolism, pH regulation, matrix metabolism, apoptosis, cytokinesis or the cell cycle. The TRX antisense oligonuclectides of the invention have increased specificity and affinity for TRX mRNA, and are resistant to degradation. The present sequence represents a poly (dT)12-18 oligonuclectide used in first strand synthesis of human thioredoxin cDNA in an example of the invention.

Revised record issued on 24-MAR-2005 : Correction to organism field

Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

Gaps ö DB 1; Length 18; 8.7e+02; 0; Indels 0.7%; Score 18; DB 100.0%; Pred. No. 8.7 tive 0; Mismatches Query Match
Best Local Similarity 100.0
Matches 18, Conservative

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2709 AAAAAAAAAAAAAA 2726 AAAAAAAAAAAAAAAA 18

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ADW10182 standard; DNA; 18 BP RESULT 1098 ADW10182/ ü

Poly (dT)12-18 oligo, SEQ ID NO:741, used in first strand cDNA synthesis. Wissenbach M; /\*tag= a /note= "Optionally and serially deleted" Westergaard M, Petersen KD, Antisense therapy; apoptosis stimulation; ss. Location/Qualifiers 10-FEB-2004; 2004US-00776934 10-FEB-2003; 2003US-0446372P. (first entry) THRUE C A. WESTERGAARD M. (PETE/) PETERSEN K D. (WISS/) WISSENBACH M. Thrue CA, (HANS/) HANSEN B. US2005014712-A1. 07-APR-2005 misc\_feature 20-JAN-2005. Synthetic. Hansen B, (WEST/) THRU/) 

New oligomeric compound for the modulation of survivin, useful for treating e.g. cancers, atherosclerosis, psoriasis, diabetic retinopathy, rheumatoid arthritis, asthma, warts, or allergic dermatitis. WPI; 2005-100663/11.

Example 6; SEQ ID NO 741; 264pp; English.

The invention relates to antisense oligomucleotides consisting of 8-50 mucleotides and/or nucleotide analogs which inhibit expression of human survivin, an inhibitor of apoptosis which is also essential for cell division and angiogenesis. The antisense oligomucleotides comprise a subsequence of 8 or more nucleotides or nucleotide analogs, wherein the subsequence of 8 or more nucleotides or nucleotides analogs, wherein the subsequence is located within a sequence selected from ADW0944-ADW09586. The oligomucleotides preferably contain a linkage group selected from a phosphate group, a phosphorothioate group or a boranophosphate group. The invention also crelates to a conjugate comprising a survivin antisense oligomucleotide of the invention and one or more non-nucleotide or non-polymucleotide of composition comprising a survivin antisense oligomucleotide, and a pharmaceutical composition comprising a survivin antisense oligomucleotide or conjugate of the invention, optionally further comprising a chemotherapeutic agent. The survivin antisense oligomucleotides, and conjugates and compositions correlating them, are useful in the treatment of cancers such as carcinoma, breast carcinoma, non-small cell lung cancer, renal cell carcinoma, bladder carcinoma, non-small cell lung cancer, renal cell carcinoma, prostatic carcinoma, prostatic carcinoma, prostatic carcinoma, cervical carcinoma, prostatic carcinoma, and kaposi's sarcoma (e.g., carcinoma, prostatic carcinoma, prostatic carcinoma, and kaposi's sarcoma (e.g., carcinoma, prostatic arcinoma, and kaposi's sarcoma, cumpanian properment of conditions such as atherosclerosis, psoriasis, diabetic certinopathy, rheumatoid arthritis, asthma, warts, and allergic certinoma proprieting apoptorial and personal edicenting and proliferation, for modulating apoptosis and for treating a disease related to abnormal angiogenesis. The survivin antisense oligonucleotides of the invention are shorter than prior art survivin antisense ö

Gaps

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Indels

Length 18;

Score 18; DB 1; Le Pred. No. 8.7e+02;

0.7%; Scor. 100.0%; Pred. No. c. 0; Mismatches

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Conservative

Local Similarity es 18; Conserv

Matches

Query Match

Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;

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The invention relates to a polynucleotide (I) for synthesizing labeled protein and having ability to increase labeling efficiency of labeling compound, where protein is produced by adding labeling compound to 3, terminal of sequence encoding target protein of gene template, where labeling compound has label portion and acceptor portion having compound capable of binding to C-terminus of label portion and translating gene template in presence of labeled compound. (I) is useful for producing a labeling protein, which involves preparing a gene template by adding (I) to the 3'-terminal of base sequence encoding the target protein, translating the gene template in the presence of the labeling compound containing acceptor portion and label portion, and obtaining protein synthesized in the translation system. The base sequence encoding the target protein either contains the termination codon or does not contain the termination codon. The labeling compound is added after the initiation of the translation. The labeled protein (LPI) is useful in a performance-analysis of a protein, which involves contacting the test substance with (LPI), and analyzing the interaction between the protein and the test substance. (I) has the ability to increase labeling efficiency of a labeling compound and thus effectively produces labeled protein. This sequence corresponds to a sequence used in the method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel polynucleotide having ability to increase labeling efficiency of labeling compound, useful for synthesizing labeled protein in presence of labeling compound.
                                                                                                                                                                                              Gaps
oligonucleotides (16-mers compared to 20-25-mers), therefore having increased specificity and affinity for survivin mRNA, and also have higher biostability and cell permeability. The present sequence represents a poly (dT)12-18 oligonucleotide used in first strand synthesis of human survivin cDNA in an example of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suzuki K;
                                                                                                                                                      DB 1; Length 18;
8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA purification; protein engineering; diagnosis; ss.
                                                                                                                Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kobayashi T,
                                                                                                                                                                100.0%; Prec. ...
                                                                                                                                                      0.7%; Score 18;
00.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein labelling method sequence #21.
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                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                          BP.
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                                                                                                                                                                                                                                                                                                                                                                    ADW86820 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                07-APR-2005 (first entry)
                                                                                                                                                  Query Match
Best Local Similarity 100.0
Matches 18, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2005-075248/08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hashimoto H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
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DNA purification; protein engineering; diagnosis; ss.

Protein labelling method sequence #18.

(first entry)

07-APR-2005

ADW86817;

ADW86817 standard; DNA; 18

RESULT 1100

ADW8681.

Suzuki K;

Kobayashi T,

Shiratori M,

Hashimoto H, Sasaki T;

Nakano H,

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18-JUN-2004; 2004WO-JP008953. 18-JUN-2003; 2003JP-00173634. (MITU ) MITSUBISHI CHEM CORP.

WO2004113530-A1. Unidentified

29-DEC-2004.

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The invention relates to a polynucleotide (I) for synthesizing labeled protein and having ability to increase labeling efficiency of labeling compound, where protein is produced by adding labeling compound to 3, cemplate of sequence encoding target protein of gene template, where labeling compound has label portion and acceptor portion having compound capable of binding to C-terminus of label portion and translating gene template in presence of labeled compound. (I) is useful for producing a labeling protein, which involves preparing a gene template in presence of labeled compound. (I) is useful for producing to the 3'-terminal of base sequence encoding the target protein, translating the gene template in the presence of the labeling compound containing acceptor portion and label portion, and obtaining protein synthesized in the translation system. The base sequence encoding the target protein either contains the termination codon or does not contain the termination codon. The labeling compound is added after the initiation of the translation. The labeled protein (LPI) is useful in a containing compound is added after the initiation of the translation. The labeled protein (LPI) is useful in a performance-analysis of a protein, which involves contacting the test cubstance with (LPI), and analyzing the interaction between the protein and the test substance with (LPI), and the shilty to increase labeling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polynucleotide having ability to increase labeling efficiency of labeling compound, useful for synthesizing labeled protein in presence of labeling compound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            efficiency of a labeling compound and thus effectively produces labeled protein. This sequence corresponds to a sequence used in the method of
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Best Local Similarity
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the invention

Screening; diagnosis; therapeutic; inflammation; antiinflammatory; cardiovascular disease; nephrotropic; endocrine disease; genitourinary disease; secreted factor; ss.

Oligo used to identify rat P00188\_D12 clone gene, SEQ ID NO: 10.

(first entry)

22-SEP-2005

AEB68153;

BP.

AEB68153 standard; DNA; 18

RESULT 1102

AEB68153

2709 AAAAAAAAAAAAAA 2726

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor protein referred as VSHK-1 useful for diagnosing or treating cancer. The invention also relates to a method for reducing the growth of a cancerous cell. VSHK-1 is useful for identifying a VSHK-1 receptor ligand and to identify a substance which modulates its signal transduction activity. The HX2004-6 DNA is useful to detect the presence of HX2004-6 in a biological sample (e.g. ductal epithelial cells from tissue chosen from pancreas, colon and breast). The invention is useful for screening drugs for the treatment of cancer. The present sequence is a reverse transcriptase (RT)-PCR primer used in the expression analysis
                                                                                                                                                                                                                                                        diagnosis; therapeutic; cancer; cytostatic; neoplasm; ss; transcriptase PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human HX2004-6 protein and a seven transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated human HX2004-6 polypeptide or isolated VSHK-1 polypeptide, useful for diagnosing or treating cancer, where VSHK-1 is also used to identify a VSHK-1 receptor ligand.
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   Gaps
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                                                                                                                                                                                                                            RT-PCR primer used in the expression analysis of HX2004-6.
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   0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
 Mismatches
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                             2709 AAAAAAAAAAAAAAA 2726
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14-MAY-1999; 99US-01144856P.
14-MAY-1999; 99US-0134112P.
13-AUG-1999; 99US-0148936P.
13-AUG-1999; 99US-0148936P.
12-MAY-2000; 2000US-00570593.
25-AUL-2000; 2000US-00570593.
25-AUL-2000; 2000US-005270593.
21-FEB-2001; 2001US-0271254P.
21-FEB-2001; 2001US-0371254P.
31-FEB-2003; 2003US-00360648.
30-OCT-2003; 2003US-005960848.
                                                                                                                                     BP.
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                                                       1 AAAAAAAAAAAAAAA
                                                                                                                                    AEA89505 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                    US2005130926-A1.
                                                                                                                                                                                                                                                          Drug screening;
RT-PCR; reverse
                                                                                                                                                                                                                                                                                                      Unidentified.
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18;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                               Screening subject for cardiac, renal or inflammatory disease caused by differential expression of polypeptide (new secreted factor) or its endogenous homologue, by measuring and determining expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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100.0%; Pred. No. 8.7e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; SEQ ID NO 10; 38pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide or its homologue.
                                                                                                                                                                                                                                                      20-SEP-2000; 2000US-00665976.
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                          Stanton LW, Kapoun AM;
                                                                                                                                                                                                                                                                                (STAN/) STANTON L W. (KAPO/) KAPOUN A M.
                                                                                                                                                                                                                                                                                                                                                     WPI; 2005-532124/54.
                                                                                                                                          Rattus norvegicus
                                                                                                                                                                    US2005158729-A1.
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Gaps

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0.7%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 8.7e+02; tive 0; Mismatches 0; Indels

Best Local Similarity 100.0 Matches 18; Conservative

Query Match

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Gaps

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0; Mismatches

18; Conservative

Matches

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The present invention provides compositions and methods for the detection of a target polynuclectide. A claimed probe set comprises: (a) a donor polymer comprising (i) a first polynuclectide binding domain complementary to a first region of a target polynuclectide flanked by first and second stem regions which hybridize in the absence of the target polynuclectide form a stem-loop or random-coil structure, and (ii) a quantum dot; and (b) an acceptor polymer comprising (i) a second plynuclectide binding domain complementary to a second region of the polynuclectide flanked by first and second stem regions which the absence of the target polynuclectide flanked by first and second stem regions which hybridize in the absence of the target polynuclectide to form a stem-loop or random-coil structure, and (ii) at least one reporter. Energy transfer occurs between the donor and the reporter(s) when the donor polymer and the reporter(s) when the donor polymer the acceptor polymer hybridize to the target polynuclectide and the cquantum dot is exposed to an exciting amount of energy. The polymers may further comprise a protein transduction domain and/or targeting signal. Preferably, at least one polymer comprises a peptide nucleic acid, a comprising a danor probe and acceptor probe. A claimed method of detecting a target polynuclectide comprises delivering at least one probe and acceptor probe. A claimed method of detecting a target polynuclectide comprises delivering at least one probe and acceptor probe hybridize to the target polynuclectide. The detectable signal is an energy polynuclectide is comprised between the quantum dot and the reporter to produce a point the target polynuclectide. The detectable signal is an energy polynuclectide or materion, deletion or insertion in the target polynuclectide, or of a certain agent polynuclectide. The detectable signal is an energet polynuclectide or method for an excit method for a certaing cells expressing a target nucleic acid, a method for sorting cells expressing a target nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New activable probe set comprising a donor polymer and an acceptor polymer, useful for in vivo gene detection or for detecting a target polynucleotide.
DNA detection; RNA detection; FRET; hybridization; probe; ss.
                                                                                                                                                 /mod_base= OTHER
/note== "OTHER= 5' Cyanine 5 fluorophore"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; SEQ ID NO 69; 147pp; English.
                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   using dual FRET molecular beacons.
                                                                                                                                                                                                                                                                                                                                                                                                          (GEOR-) GEORGIA TECH RES CORP.
                                                                                                                                                                                                                                                                                                                                          21-JAN-2004; 2004US-0538381P.
21-JAN-2004; 2004US-0538382P.
                                                                                                                                                                                                                                                                                                  21-JAN-2005; 2005WO-US001771
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                                                                                                                                                                                                                 #02005071115-A1
                                                                                                         modified base
                                                                                                                                                                                                                                                           04-AUG-2005
                                              Synthetic
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present sequence is that of a fluorescently labeled probe, which was used as a control in a fluorescence in situ hybridization (FISH) assay in an example from the invention describing mRNA detection in living cells
                                                                                                                                                                                                                          Sequence 18 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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Length 18;

0.7%; Score 18; DB 1; I 100.0%; Pred. No. 8.7e+02;

Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for augmenting T-helper 1 cells (Th1)-like immune activation in a subject. The method comprises administrating an immunostimulatory nucleic acid (I) to induce Th1-like immune activation; and administrating a cyclooxygenase inhibitor (II) to inhibit prostaglandin expression, is new. The present sequence is one such immunostimulatory nucleic acid. (I) is useful for treating nonallergic inflammatory diseases such as psoriatis, inflammatory bowel diseases (Crohn's disease and ulcerative colitis), eczema, allergic
                                                                                                                                                                                                                                                 Immunostimulant; Antiinflammatory; Antipsoriatic; Gastrointestinal-Gen.; Antiulcer; Dermatological; Antiallergic; helper T-lymphocyte; mimune stimulation; inflammation; psoriasis; inflammatory bowel disease; crohns disease; ulcerative colltis; eczema; skin allergy; contact dermatitis; ss; phosphorothioate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Use of an immunostimulatory nucleic acid and a cyclooxygenase inhibitor to augment T-helperl cells like immune activation and to treat non-allergic inflammatory diseases, e.g. psoriasis and Crohn's disease.
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100.0%; Pred. No. 8.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                  Immunostimulatory oligonucleotide, SEQ ID 882.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 882; 58pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            contact dermatitis or latex dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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2709 AAAAAAAAAAAAAAA 2726
                                                                                                                    AED75673 standard; DNA; 18 BP
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29-MAR-2002; 2002US-00112653.
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                                  1 AAAAAAAAAAAAAAAA
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Best Local Similarity 100.
Matches 18; Conservative
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modified_base
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                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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09-MAR-2006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                absence of a single target molecule or target analyte (e.g. nucleic acid, protein, lipid, bacterium). The method involves contacting sample with one or more types of nanoparticle having target binding complements and detecting any light scattering complex formed. The nanoparticle probe complexes comprise two or more probes bound to a specific target analyte. The present sequence is a T20 diluent which is used in the preparation of a shown on page 21 in example 17 of the specification. This sequence differs from the SEQ ID NO: 26 shown on page 21 in example 17 of the specification. This sequence in page 25 of the specification (see ABD67955).
                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting target (e.g. nucleic acid, protein, lipid, bacterium) in sample, comprises contacting sample with one or more types of nanoparticle having target binding complements, and detecting any light scattering complex formed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides a method for detecting the presence or
                                                                                  T20 diluent SEQ ID: 26 #2 used to prepare aptamer-coated gold probes.
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                                                                                                        Analyte detection; DNA detection; protein detection; ss.
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                                                                                                                                                                                       /mod_base= OTHER
/note= "OTHER= Linked to a steroid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                Bao YP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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100.0%; Pre-
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                   ВЪ.
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29-AUG-2003; 2003US-0499034P.
04-NOV-2003; 2003US-0517450P.
03-NAY-2004; 2004US-0567874P.
27-MAY-2004; 2004US-00854848.
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                  AED67970 standard; DNA; 18
                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                Storhoff JJ, Lucas A,
                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-784662/80.
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Best Local Similarity
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                                                                                                                                                     Key
modified_base
                                                             12-JAN-2006
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                                       AED67970;
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AEF26309 standard; DNA; 18

RESULT 1106

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AEF26309

AEF26309/ ID AEF2 XX AC AEF2

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                                                                                    diagnosis; tumor; neoplasm; cytostatic; prognosis; expression;
DNA amplification; survivin; reverse transcriptase-PCR; RT-PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting survivin expression in tested specimen, useful in detecting early stage of tumor comprises the use of PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                            Early tumor detection method associated RT-PCR primer, SEQ ID No:7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; SEQ ID NO 7; 22pp; Chinese.
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(first entry)
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nes 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ye S, Luo B;
                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                            (YESS/) YE S.
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93JP-00112515.
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                                                                                                                                  AAQ75553 standard; DNA; 19
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                                                                                                                                                                                  (first entry)
Query Match
Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-018287/03
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Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
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                                                                                                                                                                                                                                                                                                                      01-NOV-1994.
                                                                                                                                                                                                                                               aggregate;
                                                                                                                                                                                                                                                                        Synthetic.
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AAQ75554/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abblied reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                 The invention relates to one kind of ARF1 gene in cotton genome and its promoter sequence. The gene and the promoter may be preponderantly expressed in the bud, flower, fiber and boll shell. The gene and the promoter may be useful in the research of the development of cotton reproductive organ, the quality improvement and the specific expression of foreign gene in cotton reproductive organ. The present sequence represents a cotton ARF1 related PCR primer.
                                                                                                                                                                                                                                                                                      Gaps
                                                                      Adenosine phosphate-ribosylation-factor 1 gene in cotton and its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis, gene expression, reverse transcription, primer, cDNA, aggregate, restriction enzyme, ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                           DB 1; Length 18;
8.7e+02;
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                                                                                                                                                                                                                                                                    100.0%; Prec...
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                                                                                                                                                                                                                                                            0.7%; Score 18;
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 (BIOT-) BIOTECHNOLOGY RES INST CAAS
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                                                                                                           Example 1; Page 4; 13pp; Chinese.
                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                                                                                                                  AAQ75552 standard; DNA; 19 BP.
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Matches 18; Conservative
                         Ren M, Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03
                                                  WPI; 2005-659717/68
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                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ75552
                                                                                     promoter
                                                                                                                                                                                                                                                              Query Match
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                         Guo S,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                            gene expression; reverse transcription; primer; cDNA; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                       Reverse transcription primer used in cDNA analysis technique.
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Length 19;
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DB 1; Ler
8.9e+02;
thes 0;
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Pred. No. 8.9e+02;
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0.7%; Score 18; DB
100.0%; Pred. No. 8.9
ive 0; Mismatches
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by digestion with restriction enzymes.
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The present invention describes a method for describing a nucleic acid probe, which comprises using chain extending enzymes to elongate probes. The method comprises: (a) treating the sample with a chain terminating caegant to prevent polynucleotide chain growth from the nucleic acid in the sample; (b) contacting the sample with the probe containing a terminus capable of elongation by a chain extending enzyme, where the complex to the nucleic acid in the sample; (c) contacting the sample with a chain extending enzyme and its substrates, which elongates the probe; and (d) detecting the alongated hybridised probe. Also cample with a chain extending the alongated hybridised probe. Also described is a method comprising: (a) treating nucleic acid molecules or modified nucleic acid chains unextendable by a non-template-dependent enzyme; (b) hybridising the treated molecules with a nucleic acid probe that includes an extendable terminus, under conditions where hybrids form; and (c) treating any hybrids formed with a non-template dependent chain elongating enzyme and its substrates, where any hybridised probe is elongating enzyme and its substrates, where any hybridised probe is acide, particularly DNA hybridisation probes. The present sequence represents a tailing reaction exemplary primer, which is used in an example from the present invention
                                                                Identifying and detecting nucleic acids, particularly DNA hybridization probes, involves employing chain extending enzymes (e.g. telomerase) to elongate probes to render them readily detectable.
                                                                                                                                                                                                  The present invention describes a method for detecting a nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 18 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
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/note= "phenoxazine"
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                                                                                                                                                              Example 1; Page 5; 10pp; English.
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/mod base= OTHER
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28-NOV-2001; 2001US-00996292.
10-DEC-2001; 2001US-00013295.
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Best Local Similarity 100.0
Matches 18; Conservative
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                       WPI; 2002-361176/39.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AACS547-C57598) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression - by amplification of mRNA followed
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25-MAY-2000; 2000US-00580358.
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(STRE/) STREIFEL J A.
                                                                                                                                                                                                                                                                                             WPI; 1995-018287/03.
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                                                                                                                                                                                                  16-APR-1993;
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ADG28486/c
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                                                                                                          The invention relates to novel nuclease-resistant oligomeric compounds. The compounds of the invention are useful as pharmaceuticals for topical administration such as transdermal patches. The oligomeric compound is resistant to enzymatic degradation. The sequences shown in ABZ75345. ABZ75399 represent the nuclease-resistant compounds of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to novel nuclease-resistant oligomeric compounds. The compounds of the invention are useful as pharmaceuticals for topical administration such as transdermal patches. The oligomeric compound is resistant to enzymatic degradation. The sequences shown in ABZ75345-
                                                                                                                                                                                                                      Gaps
                                                    Nuclease-resistant oligomeric compound useful as pharmaceuticals for topical administration such as transdermal patches.
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                                                                                                                                                                         Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
           Rajeev KG;
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           Prakash TP,
                                                                                      Disclosure; Page 234; 234pp; English
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28-NOV-2001; 2001US-00996292.
10-DEC-2001; 2001US-00013295.
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            Manoharan M, Maier MA,
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les 18; Conserv
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modified_base
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28-NOV-2001;
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AB275399 represent the nuclease-resistant compounds of the invention
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0.7%; Score 18; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 8.9e+02;
Matches 18; Conservative 0; Mismatches 0; Indels
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                                                     Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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100.0%; Pred. No. 8.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG85220 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG28486 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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The invention relates to an isolated nucleic acid molecule (I), comprising a nucleic acid sequence chosen from 75 Nicotiana-derived cytochrome P450 enzyme fragment sequences. (I) is useful for producing a transgenic tobacco plant, which involves operably linking (I) with a promoter functional in the plant to create a plant transformation vector, and transforming the plant with the plant transformation vector, and transforming the plant with the plant transformation vector, selecting a plant cell transformed with the transformation vector, are resenctating a plant from the selected plant cell. The nucleic acid molecule is in an antisense orientation, sense orientation or is in a RNA interference orientation. The present sequence represents a PCR primer used to clone DNA encoding tobacco cytochrome P450 enzyme fragments of
                                                                                                                                                                                                                                                                                                                                                                                                                                    An isolated nucleic acid molecule, comprising nucleic acid sequence of Nicotiana derived cytochrome P450 enzyme fragments, useful for producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplified RNA preparation; nervous system disorder;
neurodegenerative disease; Parkinson's disease; Alzheimer's disease;
multiple sclerosis; psychiatric disorder; schizophrenia;
affective disorder; manic depression; lack of appetite control;
attention deficit disorder; cancer cell detection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          represents a T7 RNA polymerase promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amplified RNA (aRNA) preparation method-related DNA sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.7%; Score 18; DB 1; Length 19;
100.0%; Pred. No. 8.9e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 73; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                                                                                     (USSM-) US SMOKELESS TOBACCO CO.
                                                                                                                                                                                                            11-JAN-2002; 2002US-0347444P.
12-MAR-2002; 2002US-0363684P.
10-JAN-2003; 2003US-00340861.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           H.
                                                                                                                                                                 12-MAR-2003; 2003US-00387346.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADT55005 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/note= "N 1
sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-DEC-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-449487/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transgenic plants.
                                                                           US2004117869-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           misc_difference
                              Nicotiana sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention
                                                                                                                     17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADT55005;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        원
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes an oligonucleotide comprising several nucleotides covalently linked together by internucleotide linkages. At least one of the nucleotides is linked to an adjacent nucleotide by 2'.5'.

The nucleotide sinkage and bears a 3'-substituent. The oligonucleotides internucleotide linkage and bears a 3'-substituent. The oligonucleotides are useful: as antisense oligonucleotides; in pharmaceutical compositions of protein e.g. organism that utilises RNA-DNA transcription or RNA-protein to protein e.g. organism that utilises RNA-DNA transcription or RNA-protein translation, bacteria, yeast, protozoa, algae and warm-blooded animals; con developing diagnostic and therapeutic agents. The modified coligonucleotide exhibits improved properties of nuclease resistance and coligonucleotide is potent antisense agent with longer cucleic acids. The oligonucleotide is potent antisense agent with longer duration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligonucleotide comprising at least one 2',5'-internucleotide linkage useful for treating organisms having disease caused by undesired
                                                                                      antibacterial; protozoacide; antialgal; fungicide; internucleotide linkage; 2',5'-internucleotide linkage; 3'-substituent; antisense; pharmaceutical; RNA-DNA transcription; RNA-protein translation; infection; diagnostic; therapeutic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          production of protein e.g. bacteria, yeast, protozoa and algae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; primer; PCR; cytochrome P450; transgenic; tobacco; plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.7%; Score 18; DB 1; Length 19;
100.0%; Pred. No. 8.9e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 54; SEQ ID NO 7; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tobacco cytochrome P450 PCR primer #14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Guinosso CJ;
                                            Modified oligonucleotide seg id 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAAA 2726
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94WO-US010131.
96US-00602862.
98US-00115043.
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                                                                                                                                                                                                                                                                                                                                                                  99US-00435806
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26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-SEP-2004 (first entry)
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watches 18; Conservative
                                                                                                                                                                                  nuclease resistance; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cook PD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-079586/08.
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                                                                                                                                                                                                                                                                         US6653458-B1
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28-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-1993;
                                                                                                                                                                                                                                                                                                                     25-NOV-2003
                                                                                                                                                                                                                            Synthetic.
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ઠે d RESULT 1116 ADO59144

Query Match

ADO59144;

**EXEXEXEXE** 

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Gaps

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New nucleic acid molecule encoding cytochrome P450 enzymes in Nicotiana,
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                                                          WPI; 2005-223176/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADZ65660;
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                cancer.
                                    Bao G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Xu D;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADZ65660/
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                                                                                                                                                                                                                                                                 The invention comprises a method of preparing amplified RNA (aRNA)

sequences present in one or more target polynucleotide that is single
stranded or made single stranded. The method involves forming double
stranded or made single stranded. The method involves forming double
stranded cDNA templates containing sequences present in the target
copple of initiating transcription via the promoter region to produce
amplified RNA containing sequences of the target polynucleotide. The
method of the invention is useful for amplifying the population of RNAs
contained formal inflating sequences of the target polynucleotide. The
stracted from formallin-fixed tissues and/or the population of RNAA
splice variants. The method is also useful for determining gene
captession in neuronal and non-neuronal cells involved in disorders of
the nervous system, such as: neurodegenerative diseases (e.g. Parkinson's
disorders (e.g. schizophrenia); and multiple sclerosis); psychiatric
cdisorders (e.g. schizophrenia); and affective disorders (e.g. manic
cdepression, lack of appetite control, and attention deficit disorder).
The method of the invention may also be used to detect cancer cells, and
copilitate diagnosis/prognosis of cancer in a patient. The present DNA
copilitate diagnosis/prognosis of cancer in a patient. The present DNA
copilitate diagnosis/prognosis of cancer in a patient. The present DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                              Preparing amplified RNA (aRNA) sequences present in single stranded or made single stranded target polynucleotide(s), useful for detecting cancer cells, comprises transcribing double stranded cDNA templates with an RNA polymerase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Molecular beacon; DNA detection; RNA detection; probe; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 18; DB 1; Length 19;
.00.0%; Pred. No. 8.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                              Enright E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Preu. ...
                                                                                                                              Ma X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAAA 2726
                                                                                                                                                                                                                                              Example 1; Fig 1; 46pp; English.
                                                                                                    (ARCT-) ARCTURUS BIOSCIENCE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADY58870 standard; DNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JUN-2003; 2003US-0482648P.
23-JUN-2004; 2004US-00874920.
                                                  19-MAR-2004; 2004WO-US008553
                                                                            21-MAR-2003; 2003US-0456825P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Erlander MG, Salunga RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                      WPI; 2004-710328/69.
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WO2004085681-A2
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                          07-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADY58870;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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The invention provides nucleic acid reporters and methods of their use.

The nucleic acid reporters include molecular beacons modified with protein transduction domains (PTDS) to facilitate translocation of the nucleic acid reporter across cellular membranes. The nucleic acid reporter across cellular membranes. The nucleic acid reporter to a specific cell, tissue, organ, citracallular region, organelle or vesicle. The molecular beacon can be used in methods for: detecting or sorting cells expressing a target nucleic acid, in a living cell; identifying modulators of gene expression; determining the effectiveness of an agent on a host cell; and detecting the transport and localization of a nucleic acid, in a living cells expressed on a specific cell; and detecting the transport and localization of a nucleic acid in a living cells; identifying colls acid-protein complex in living cells. The present sequence is of a polyh probe. This was used as a negative control in a fluorescence in situ hybridization (Fish) assay targeting GAPDH mRNA in a comparison of this traditional method of detection with the method of the invention.
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                                                                                                                                                                                                                                      New molecular beacon operably linked to a protein transduction domain, useful in preparing a pharmaceutical composition for treating e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytochrome p450; 88; secondary metabolite; ethylene; senescence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 0.7%; Score 18; DB 1; Length 19; Local Similarity 100.0%; Pred. No. 8.9e+02; les 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 18 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oilgo d(T) PCR primer for amplifying p450 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                        Example 9; SEQ ID NO 22; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nornicotine; transgenic plant; primer.
                                                                                    Kim 67;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAAA 2726
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(GEOR-) GEORGIA TECH RES CORP.
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29-APR-2004; 2004US-0566235P.
03-SEP-2004; 2004US-00934944.
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ADZ66018 standard; DNA; 19 BP.
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                                                                                                                                                                                                               15-OCT-2004; 2004WO-US034218
                                                                                                                                                    14-JUL-2005 (first entry)
                                                                                                                                                                                                                                                             WPI; 2005-315709/32.
                                                                                      Best Local Similarity
                                                                                                                                                                               Nicotiana tabacum.
Synthetic.
                                                                                                                                                                                             WO2005038018-A2.
                                                                                                                                                                                                                                                                                nornicotine
                                                                                                                                                                                                      28-APR-2005
                                                                                                                                            ADZ66018;
                                                                                 Query Match
                                                                                                                          RESULT 1120
                                                                                          Matches
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                                                                                                                               ADZ66018,
#X8X555555555555X8
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nucleic acid molecule, a method of increasing or decreasing nornicotine levels in a plant by operably linking the nucleic acid molecule with a promoter functional in the plant, a tobacco product having reduced amounts of nornicotine levels, the tobacco product comprising tobacco from the plant, a tobacco leaf having reduced amounts of nornicotine levels and a method of isolating a gene from a plant using the isolated nucleic acid. In producing a transgenic plant, the plant has reduced levels of nornicotine. The tobacco product is selected from cigarettes, cigars, pipe tobacco, snuff, chewing tobacco, products blended with the tobacco product and their mixtures. The nucleic acid molecule is useful for altering plant phenotypes, thus producing a transgenic plant having reduced levels of nornicotine. This sequence represents a PCR primer used to amplify CDNA encoding a tobacco cytochrome P450 enzyme of the Score 18; DB 1; Length 19; Pred. No. 8.9e+02; 0; Indels Oligonucleotide used for modulating gene transcription. Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other; Cook Z, Fang Y, Feldmann K, Kiegle EA, Kwok S, Pennell R, Schneeberger R, Wu C; 100.0%; Pred. ... Disclosure; SEQ ID NO 2; 210pp; English. cell or transformed host organism. 2709 AAAAAAAAAAAAAAA 2726 23-SEP-2003; 2003US-0505689P. 14-0CT-2003; 2003US-0511460P. 06-NOV-2003; 2003US-0518075P. 04-DEC-2003; 2003US-0527611P. 12-DEC-2003; 2003US-05293S2P. 13-FEB-2004; 2004US-0549771P. AAAAAAAAAAAAAA 1 BP 23-SEP-2004; 2004US-00950321 0.7%; AED19814 standard; DNA; 19 (first entry) 18; Conservative Similarity WPI; 2005-664198/68. Transcription; ss. (CERE-) CERES INC. US2005223422-A1. Unidentified 01-DEC-2005 06-OCT-2005. AED19814; 18 Query Match Local RESULT 1121 Matches AED19814/C
XX
AC
AED1:
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AED1:
AED1 ò 원 ö The invention relates to an isolated nucleic acid molecule (I) from Nicotiana, where the nucleic acid molecule comprising any of the 59 nucleic acid sequences of SEQ ID NOS: 299-357 (NOTE:The claims refer to SEQ ID NOS 299-357 as nucleic acids but these sequences (apart from SEQ ID NO 356) are all proteins and appear as ADZ65402-ADZ65460. The nucleic acids of the invention encode cytochrome p450 enzymes whose expression is induced by ethylene and/or plant senescence. Also included are a transgenic plant comprising (I), a method of producing a transgenic plant, a method of selecting a plant containing a nucleic acid molecule where the plant is analyzed for the presence of nucleic acid sequence of Nicoliana, encoding a protein. The invention also relates to a transgenic plant comprising the nucleic acid molecule, a method of producing a transgenic plant comprising operably linking the nucleic acid molecule with a promoter functional in the plant to create a plant transformational vector, transforming the plant with the plant transformational vector, selecting a plant cell transformed with the transformation vector and regenerating a transformation plant from the transformation vector and regenerating a transformation plant from the transformed plant cell, a method of selecting a plant containing a New isolated nucleic acid molecule from Nicotiana, useful for altering plant phenotypes, thus producing a transgenic plant having reduced levels Gaps .; 0 invention relates to an isolated nucleic acid molecule from useful in developing tobacco plants with altered phenotypes. Score 18; DB 1; Length 19; Pred. No. 8.9e+02; 0; Indels Enzyme engineering; cytochrome P450; PCR; ss; primer. Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other; Tobacco cytochrome P450 enzyme cDNA PCR primer #6. 100.0%; Pred. nc. Disclosure; Fig 152; 226pp; English. Disclosure; Fig 153; 203pp; English. 2726 (USSM-) US SMOKELESS TOBACCO CO. 18 AAAAAAAAAAAAAAA 1 2709 AAAAAAAAAAAAAA 18; Conservative

The invention relates to a nucleic acid molecule or its complement sequence capable of modulating transcription. The nucleic acid molecule of the invention is useful for transcription of polynucleotides in a host cell or transformed host organism. The present sequence is an oligonucleotide used for modulating gene transcription. New isolated nucleic acid molecule capable of modulating transcription, or its complement, useful for transcription of polynucleotides in a host

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The invention relates to an isolated nucleic acid molecule capable of modulating transcription, where the nucleic acid molecule shows at least combulating transcription, where the nucleic acid molecule shows at least specification or its complement. The invention also relates to a vector construct comprising a first nucleic acid capable of modulating transcription, where the nucleic acid molecule shows at least 80% transcription, where the nucleic acid molecules are heterothy to one of the promoter sequences given in the sequence identity to one of the promoter sequences given in the specification, and a second nucleic acid molecules are heterologous to each other and are operably linked together, a host cell comprising the nucleic acid molecule is flanked by an exogenous concerning the comprising the vector construct. The first comprising the vector construct. The first comprising the vector construct. The first comprision the vector construct. The first comprision in the specification, where the plant cell is regenerated concours in the specification, where the plant cell is regenerated into a plant. The nucleic acid molecules, which are promoter sequences, cor comprision in the specification, where the plant cell is regenerated into a plant. The nucleic acid molecules, which are promoter sequences, and promoter control elements are useful for modulating transcription of a desired polymucleotide in plantes. This sequence represents a synthetic continue to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated promoter sequences and promoter control elements, useful for modulating transcription of a desired polynucleotide in plants.
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Pred. No. 8.9e+02;
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100.0%; Pred. No. c...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wu C;
                                                                                                 Transcription; vector; primer; ss.
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04-DEC-2003; 2003US-0527611P.
13-FEB-2004; 2004US-0544771P.
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                   29-DEC-2005 (first entry)
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Best Local Similarity 100.
Matches 18; Conservative
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                                                            Synthetic primer #2.
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                                                                                                                                                                                   US2005246785-A1.
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Pennell R,
                                                                                                                                                                                                                          03-NOV-2005
                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleotide sequences, useful modifying plant characteristics or for modulating and manipulating growth, development, and biochemistry of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                      0.7%; Score 18; DB 1; Length 19;
100.0%; Pred. No. 8.9e+02;
tive 0; Mismatches 0; Indels
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Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;
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Wu C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alexandrov N, Zhihong C, Fang Y,
Lu Y, Penell R, Schneeberger R,
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                                                                                                                                                                                                                                                                        AED21473 standard; DNA; 19 BP.
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                                                                                                                                                   (first entry)
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                                                                                  Matches 18; Conservative
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Matches 18; Conserv
                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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RESULT 1123 AED60796/c

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CDNAs of the invention. 

Degenerate PCR primer for cloning cytochrome p450 cDNA, SEQ ID NO: 2260.

Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;

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Plant breeding; plant; 8s; PCR; transgenic plant; senescence; primer.

Synthetic.

WO2005111217-A2.

24-NOV-2005.

2004US-0566235P. 27-APR-2005; 2005WO-US014803 29-APR-2004;

2004US-0607357P. 2004WO-US034065. 2004WO-US034218. 2005US-0646764P. 24-MAR-2005; 2005US-0665097P. 24-MAR-2005; 2005US-0665451P. 2004US-00943507 15-OCT-2004; 25-JAN-2005; 15-OCT-2004; 03-SEP-2004; 17-SEP-2004;

(USSM-) US SMOKELESS TOBACCO CO

Xu D, Nielsen MT;

WPI; 2005-786788/80

demethylase gene comprises crossing a first tobacco plant with a second tobacco plant and germinating the collected seed of an Fl progeny plant Producing a tobacco plant having decreased expression of a nicotine

Disclosure; SEQ ID NO 2260; 641pp; English.

The invention relates to a breeding method for producing a tobacco plant with reduced expression of a nicotine demethylase gene expression af first tobacco plant with at least one phenotypic trait to produce a first tobacco plant with at least one phenotypic trait to produce a first tobacco plant with at least one phenotypic trait to produce a tobacco plant having reduced expression of a nicotine demethylase deficiency trait into a tobacco plant, producing a tobacco seed, deficiency trait into a tobacco plant, producing a tobacco plant in a tobacco beained from the tobacco plant conting a modified attribute, a method for the invention, producing a tobacco plant avoing a modified attribute, a method of preeding an attribute into a tobacco plant, producing a tobacco plant paying a modified attribute of method of preeding an attribute into a tobacco plant producing a tobacco plant producing a tobacco plant producing a tobacco plant producing a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or developing a tobacco plant baving a modified attribute or sequence given in the specification (the nucleic acids comprise isolated corrected by the modified attribute or senescence induced from a germinated seed of the plant, reducing substantially identical to a michigan or enzymatic activity of a constitutive, or an ethylene or senescence induced tobacco polypeptide in a plant cell, the expression or enzymatic activity of a constitutive, or an ethylene or senescence induced tobacco polypeptide in a plant produced to senescence induced toba

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                                                                                                                                                                                                                                                                                                                                    New nucleic acid molecule, useful in producing transgenic plants for use as models for modifying plant characteristics e.g. increase in plant height, number or size of leaves, or wood products.
                   Gaps
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                                                                                                                                                                                                                                                                                            Fang Y, Kwok S,
 Length 19;
                 0; Indels
0.7%; Score 18; DB 1; Le
100.0%; Pred. No. 8.9e+02;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                            Schneeberger R,
                                  2709 AAAAAAAAAAAAAAA 2726
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                                                   18 AAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                          12-DEC-2003; 2003US-0529352P.
                                                                                                                                                                    ss; primer; transgenic plant
                                                                                            AEF26614 standard; DNA; 19
                                                                                                                                 (first entry)
                  18; Conservative
                                                                                                                                                                                                                                                                                           Pennell R, Okamuro J,
                                                                                                                                                                                                                                                                                                     Donson J,
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                          (CERS-) CERS INC.
                                                                                                                                                  Oligo dTV primer.
                                                                                                                                                                                                     JS2006015970-A1.
                                                                                                                                 23-MAR-2006
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                                                                                                                                                                                                                                                                                                   Kiegle EA,
                                                                                                                                                                                     Synthetic.
                                                                                                                AEF26614;
                                                                              RESULT 1125
                  Matches
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Arabidopsis thaliana and the polypeptides encoded by them. The patentees also claim a vector construct containing such nucleic acid molecules, and a laso claim a vector construct containing such nucleic acid molecules, and a host cell transformed; a method for detecting a nucleic acid in a sample; and a plant, plant cell, plant material or seed of a plant which comprises the nucleic acid molecule which is exogenous or heterologous to the plant or plant cell. The vector construct comprises a first nucleic acid having a regulatory sequence capable of causing transcription and/or translation in a plant and a second nucleic acid having the sequence of the isolated nucleic acid molecule. The first and second nucleic acids are operably linked and where the second nucleic acid is heterologous to any element in the vector construct. The isolated nucleic acid molecules are useful in producing transgenic plantes for use as models for modifying plant characteristics, e.g. increase in plant height, number or size of leaves, or wood products. The present sequence is that of a primer used in generation of labeled probes from first-stand cDNA.

Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;

Gaps ; 0 0.7%; Score 18; DB 1; Length 19; 100.0%; Pred. No. 8.9e+02; Indels 100.0%; Pred. ... 18; Conservative Local Similarity Query Match Matches

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2709 AAAAAAAAAAAAAA 2726 18 AAAAAAAAAAAAAAAAA

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Plant breeding; crop improvement; secondary metabolite; genetic marker; reverse transcriptase PCR; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                            New breeding method, useful for producing a tobacco plant having decreased expression of a nicotine demethylase gene comprises crossing germinating F1 progeny seed to produce the tobacco plant.
                                                                    Oligo(dT) reverse RT-PCR primer, SEQ ID NO:2260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 2260; 511pp; English
                                                                                                                                                                                                                                                                                                                                                                          (USSM-) US SMOKELESS TOBACCO CO
                                                                                                                                                                                                                                                                                                                      2004WO-US034218.
2005US-0646764P.
2005US-0665097P.
2005US-0665451P.
                 AEF99207 standard; DNA; 19 BP.
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2003US-00340861.
2003US-00387346.
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2003US-0503989P.
2003US-00686947.
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2004US-00934944.
2004US-0607357P.
2004US-00943507.
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                                                    (first entry)
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                                                                                                                                 US2006041949-A1
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16-OCT-2003;
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12-MAR-2002
                                                                                                                                                   23-FEB-2006
                                                                                                                 Synthetic.
                                   AEF99207;
RESULT 1126
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The invention relates to a breeding method for producing a tobacco plant having decreased expression of a nicotine demethylase gene. The method involves crossing a first tobacco plant having variant nicotine demethylase gene expression with a second tobacco plant containing at least one phenotypic trait (e.g., disease resistance, high yield etc.) to produce an Pl progeny plant; collecting the seed of the Pl progeny; and germinating the seed to produce a tobacco plant having decreased nicotine demethylase expression. The invention also relates to a tobacco plant or its components produced using the method of the invention; a tobacco product produced using the method of the invention; a tobacco product produced from such plants; and a method of breeding a tobacco product produced from such plants; and a method of breeding a tobacco product produced from such plants; and a method of breeding a tobacco product produced from such plants; and a method of breeding a tobacco product produced expression of a plant with a modified attribute comprising variant expression of a cytochrome P450 polynucleotide (including the nicotine demethylase gene). The invention further relates to isolated constitutive, ethylene-induced or senescence-induced genetic markers comprising tobacco cytochrome P450 concleded increasing the expression or activity of polypeptides for reducing or increasing the expression or activity of polypeptides encoded by these nucleic acid molecules; expression vectors, plants or

plant components comprising one of these polynucleotides; and tobacco products produced from such plants. The methods of the invention are useful for the breeding) (especially marker sesisted breeding) of tobacco cytochrome P450 expression or activity. Altered expression or with altered cytochrome P450 expression or activity. Altered expression of such a crange in the composition of secondary cytochrome P450 expression or activity. Altered expression of such cytochrome P450 expression or activity. Altered expression of such cytochrome P450 expression or activity. Altered expression of such cytochrome P450 expression or activity. Altered expression of such cytochrome the flavor or aroma of plant products. They may also affect cytochrome the flavor or aroma of plant products. They may also affect herbicide tolerance, resistance to disease or insects, quality factors correlated to undesirable constituents, structural traits, disher content. The invention in which the expression of extochrome P450 genes is of the invention in which the expression of extochrome P450 genes is caltered may have desirable traits such as altered levels of nornicotine or N-nitrosonornicotine. Such plants can be used in the production of tobacco products such as moist or dry snuff, chewing tobacco, cigarettes, cigarillas, pipe tobacco, bidis or smokeless tobacco products controlled to activity of tobacco cytochrome P450 charactivity from the US patent is expression of controlled to also available in electronic format directly from the US patent of fice at also available in electronic format directly from the US patent tief controlled to also available to the format directly from the US patent of fice at segdata.uspto.gov/seguence.html?DocID=20060041949 %XGGGGGGGGGGGGGGGGGGGGGG

Sequence 19 BP; 0 A; 0 C; 0 G; 18 T; 0 U; 1 Other;

Gaps . 0 Length 19; 0; Indels Score 18; DB 1; Lo Pred. No. 8.9e+02; 100.0%; Pred. ... 0.7%; Local Similarity 100. hes 18; Conservative Query Match Matches

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2709 AAAAAAAAAAAAAAAA 2726 18 AAAAAAAAAAAAAA 1 g ð

BP. AAQ75586 standard; DNA; 20 (first entry) 04-AUG-1995 AAQ75586; RESULT 1127 AAQ75586, 

Reverse transcription primer used in cDNA analysis technique.

Analysis, gene expression, reverse transcription, primer, cDNA, aggregate, restriction enzyme, ss.

Synthetic.

JP06303997-A. 01-NOV-1994. 93JP-00112515 16-APR-1993; (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.

93JP-00112515.

16-APR-1993;

WPI; 1995-018287/03.

Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.

Disclosure; Page 5; 11pp; Japanese.

븅탕 A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily

7 10:41:34 2006

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Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                             ch 0.7%; Score 18; DB 1; Length 20; Similarity 100.0%; Pred. No. 9.1e+02; 18; Conservative 0; Mismatches 0; Indels
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate The
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04-AUG-1995

AAQ75582

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RESULT 1131

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01-NOV-1994

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93JP-00112515.

16-APR-1993;

01-NOV-1994.

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Human oligonucleotide sequence
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     ABZ88694 standard; DNA; 20
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          ABZ88694;
RESULT 1134
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthemmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds

Homo sapiens

31-OCT-2002,

WO200285308-A2.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; JW, Nyce JW Miller

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 3936; 872pp; English.

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The invention relates to a novel pharmaceutical composition, which has a
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                                                                                                                                 Sequence 20 BP; 17 A; 0 C; 0 G; 3 T; 0 U; 0 Other;
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0.7%; Score 18; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 9.1e+02;
Matches 18; Conservative 0; Mismatches 0;
                                                                                                                    at ftp.wipo.int/pub/published_pct_sequences
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2725 3 TAAAAAAAAAAAAA 20 2708 TAAAAAAAAAAAAAA 셤 ઠે

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BP
   ADH67348 standard; DNA; 20
           (first entry)
           25-MAR-2004
       ADH67348;
RESULT 1135
 ADH67348,
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inflammation, tumour formation, diabetes, obesity; cardiovascular disorder, hyperlipidaemia, Cushing's syndrome, human, ss; phosphorothioate backbone, 2'-methoxyethyl; 2'-MOE. antisense oligonucleotide; glucocorticoid receptor; infection;

Human glucocorticoid receptor-specific antisense oligonucleotide #4182.

Ното варіепв.

WO2003099215-A2

04-DEC-2003.

20-MAY-2003; 2003WO-US016084.

20-MAY-2002; 2002US-0381857P

(PHAA ) PHARMACIA CORP.

Nalseth AE;

Crosby SD,

WPI; 2004-035034/03.

New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.

Claim 4; SEQ ID NO 4182; 985pp; English.

The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense

ADK74688 standard; DNA; 20 BP.

RESULT 1137

ADK74688;

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oligonuclectides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleoride that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human glucocorticoid receptor-specific antisense oligonucleotide #4235.
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                                                                                                                                                      0.7%; Score 18; DB 1; Length 20; 100.0%; Pred. No. 9.1e+02; tive 0; Mismatches 0; Indels
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                                                                                                                Sequence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 4235; 985pp; English.
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                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                        Query Match
Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-035034/03.
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Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003099215-A2.
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathy, trigeminal neuropathy, arthritic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder; including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with the annay and a decoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                 Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps.
                                                                                                                 Chimeric phosphorothioate oligonucleotide to target Navl.3 #2022.
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Pred. No. 9.1e+02;
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:ive 0; Mismatches
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es 18; Conservative
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                                                                                                                                                                                                                                                            WO2004016754-A2.
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                                                                                    20-MAY-2004
                                                                                                                                                                                                                                                                                              26-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Roberds SL;
                                                                                                                                                                                                                           Synthetic.
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ADK74367
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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralism, chronic pain, lower back pain, diabetic neuropathy, trigeninal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present but not limited a chimeric phosphorothioate oligonucleotide with human Nav1.3 expression, the oligonucleotides inhibition of different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure
                                    Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
Chimeric phosphorothioate oligonucleotide to target Navl.3 #1701.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 20 BP; 0 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 1701; 417pp; English.
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0; M
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                                                                                                                                                                                                                                14-AUG-2003; 2003WO-US025465
                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disorder, or ataxia.
                                                                                                                                                    WO2004016754-A2
                                                                                                                                                                                            26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                   Roberds SL;
                                                                                                                Synthetic
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The present invention relates to a method (M1) for labeling double stranded nucleic acid for efficient detection of DNA or RNA. The method comprises using a carbodismide derivative for labeling one or more of thymine, uracil and guanine, which exists in the mismatch region of the double stranded nucleic acid or its vicinity, or unstable region of the hydrogen bond of the double stranded nucleic acid. (M1) is useful for labeling double stranded or single stranded nucleic acid or detecting single nucleotide polymorphisms. The present sequence was used to illustrate the method of the invention.

Labeling double stranded nucleic acid, involves utilizing carbodiimide derivative for labeling thymine, uracil and guanine, which exists in mismatch region of nucleic acid or unstable region of hydrogen bond of nucleic acid.

Example 1; Page 24; 40pp; Japanese.

<u>ы</u>

Tabata

Mukumoto K,

F

Fakenaka S, Nojima WPI; 2005-685344/71.

(TAKE/) TAKENAKA S. 18-MAR-2004;

18-MAR-2004; 2004JP-00078900.

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Gaps

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0; Indels

0.7%; Score 18; DB 1; Length 20; 100.0%; Pred. No. 9.1e+02;

100.0%; Prec. ....

Local Similarity 100. nes 18; Conservative

Best Loca Matches

Query Match

2708 TAAAAAAAAAAAAA 2725

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TAAAAAAAAAAAAAAA

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BP.

AAQ75702 standard; DNA; 21

RESULT 1140

AAQ75702;

Gaps

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Score 18; DB 1; Length 20; Pred. No. 9.1e+02; 0; Mismatches 0; Indels

Query Match 0.7° Best Local Similarity 100. Matches 18; Conservative

2709 AAAAAAAAAAAAAAAA 2726

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AAAAAAAAAAAAA 3

20

standard; DNA; 20

AED13298 AED13298;

RESULT 1139

Sequence 20 BP; 17 A; 0 C; 0 G; 3 T; 0 U; 0 Other;

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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)
                                                                                       Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                              (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                           93JP-00112515
                                      (first entry)
                                                                                                                                                                                                                                                                                        WPI; 1995-018287/03
                                                                                                                                                         JP06303997-A
                                                                                                                                                                                                          16-APR-1993;
                                    04-AUG-1995
                                                                                                                                                                                                                                      16-APR-1993;
                                                                                                                                                                                   01-NOV-1994.
                                                                                                                               Synthetic.
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Oligonucleotide ODN6 used to illustrate nucleic acid labeling method.

(first entry)

01-DEC-2005

DNA detection; RNA detection; SNP detection;

JP2005265617-A.

Synthetic.

29-SEP-2005

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gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GRNESEG files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             electionuresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                           Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                     Reverse transcription primer used in cDNA analysis technique.
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100.0%; Pred. No. 9.3e+02;
vative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 7; 11pp; Japanese.
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                                   04-AUG-1995 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-018287/03.
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nes 18; Conserv
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aggregate;
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                                                                                                                                                                        Synthetic.
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 AAQ75671;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The
                                      Analysis of cDNA and gene expression - by amplification of mRNA followed
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Matches 18; Conservative
WPI; 1995-018287/03.
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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18; Conservative
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                                                                                                                                                                                                                                                                                             - by amplification of mRNA followed
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                                        transcription primer used in cDNA analysis technique.
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Pred. No. 9.3e+02;
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                                                                                restriction enzyme; ss
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AAQ75690 standard; DNA; 21 BP
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Matches 18; Conservative
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                                                               Analysis; gene
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Pred. No. 9.3e+02;
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                                                                   Disclosure; Page 7; 11pp; Japanese.
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Matches 18; Conservative
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ7547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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100.0%; Pred. No. 9.3e+02;
tive 0; Mismatches 0; Indels
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Best Local Similarity
Matches 18; Conserv
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electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 9.3e+02;
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by digestion with restriction enzymes.
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                           expression; reverse transcription; primer; cDNA;
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         Reverse transcription primer used in cDNA analysis technique.
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iive 0; Mismatches 0;
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                                       restriction enzyme; ss
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Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                A method for the analysis of cDNA comprises (a) preparing an aggregate c double-stranded cDNAs by using an aggregate of mRNAs and a plural type c labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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100.0%; Pred. No. 9.3e+02;
ive 0; Mismatches 0; Indels
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by digestion with restriction enzymes.
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                                                                  Disclosure; Page 7; 11pp; Japanese.
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Matches 18; Conserv
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                   double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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100.0%; Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type o labelled reverse transcription primers (GENESEQ files AAQ75547075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                             Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.7%; Score 18; DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. No. 9.3 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2708 TAAAAAAAAAAAAA 2725
                                                                                             BP
18 TAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                     93JP-00112515
                                                                             683/c
AAQ75683 standard; DNA; 21
                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-018287/03.
                                                                                                                                                                                                                                           Analysis; gene
                                                                                                                                                                 04-AUG-1995
                                                                                                                                                                                                                                                                                                                                    JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-1993;
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01-NOV-1994

Synthetic.

AAQ75683;

This invention describes a novel method for sorting single-stranded mucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)

JP06303997-A

Synthetic

BXZXXXXXXXXXXXXX

01-NOV-1994

04-AUG-1995

AAQ75701;

RESULT 1167

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Gaps

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of of

comprise variable and non-variable regions. The capture oligonuclectides have a S'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of single stranded nucleic acids. The capture of comprises a sequence of 10-50, particularly 15-25, T residues. The capture conjugence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting used (i) for analysis of patterns, especially in mucosal, additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and confined and provides rapid, inexpensive and reproducible capture probes in pools of mucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual comparison of nucleic acid pools, not individual capture probes used in the method of the invention. Sequence 21 BP; 0 A; 3 C; 0 G; 18 T; 0 U; 0 Other; 

Gaps ö 0.7%; Score 18; DB 1; Length 21; 100.0%; Pred. No. 9.3e+02; tive 0; Mismatches 0; Indels Query Match Best Local Similarity 100.C Matches 18; Conservative

2709 AAAAAAAAAAAAAAA 2726 

ઠે 셤 RESULT 1169

ADK01307 standard; DNA; 21 BP. ADK01307 

06-MAY-2004 (first entry) ADK01307;

Rat DNA microarray capture oligonucleotide #27.

se; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

Dieck HT, Hoppe H; DEGS ) DEGUSSA BIOACTIVES GMBH. Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded racids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable enchor region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow that comprises are particularly locked nucleic acids (LNA) and the anchor region comprises as sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues by interaction with streptavidin. The matrix is of plastic, caramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food mixtures. The method provides rapid inexpensive and reproducible conductines. The method provides rapid inexpensive and reproducible conductines or supplements, especially minerals, race elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and matrix anging of the complete pattern of all nucleic acid sol in a cell, and one is based on comparison of uniterication of unitered acid in a cell, and one and enchance matrix miniaturism in an enchance in the nucleic acid conduction is nonembrane. genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.

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Sequence 21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;

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Gaps .. 0 0.7%; Score 18; DB 1; Length 21; 100.0%; Pred. No. 9.3e+02; 0; Indels Query Match
Best Local Similarity 100.0%; Fred. No. 9.3

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2709 AAAAAAAAAAAAAAA 2726 AAAAAAAAAAAAAAA 18

ð g RESULT 1170

ADK01306 standard; DNA; 21 BP. ADK01306; 

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #26.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

DEGS ) DEGUSSA BIOACTIVES GMBH.

Ξ Hoppe Dieck HT, Boekenkamp D,

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture of reading out, where the nucleic acids are selectively bound using capture of each and immobilised on the surface of a solid matrix and (b) comprise variable anchor region. The capture oligonuclectides of the anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nuclectides to allow binding of particular sorts of single stranded nucleic acids. The capture of some comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, reshin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is quasilocal, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially, in mucosal, chair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic caids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible caid in a cell, and callows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid in a cell, and canner matrix miniarive in anniarion is not indial-acid acid in a cell, and canner matrix miniarive in a nucleic acid acid and an acceptant of an acceptance of acceptance 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention.
                                                            Example; Page 5; 8pp; German.
   and constant regions.
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## Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;

; 0 0.7%; Score 18; DB 1; Length 21; 100.0%; Pred. No. 9.3e+02; 0; Indels Local Similarity 100.0%; Pred. No. 9.3 nes 18; Conservative 0; Mismatches Query Match Best Loca Matches

## 2709 AAAAAAAAAAAAAA 2726

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ADK01305 standard; DNA; 21 BP. ADK01305; RESULT 1171 

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #25.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH

Boekenkamp D, Dieck HT, Hoppe H;

WPI; 2003-714082/68.

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture of gentuse oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, computation is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, eag food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible captow imaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual in the complete pattern of all pools of the properior of acid in a passed on comparison of nucl capture probes used in the method of the invention.

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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Gaps ö 0.7%; Score 18; DB 1; Length 21; 00.0%; Pred. No. 9.3e+02; 0; Indels 100.0%; Preq. w... 18; Conservative Best Local Similarity Query Match Matches

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ð 셤 RESULT 1172 ADK01308/

ADK01308 standard; DNA; 21 06-MAY-2004 (first entry) ADK01308; 

BP.

Rat DNA microarray capture oligonucleotide #28.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794

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Rat DNA microarray capture oligonucleotide #41.
                                                                                                              DE10208794-A1
                                                                                                           Rattus sp.
                                                                                            ADK01321;
                                                                                      RESULT 1173
                                                                                        ADK01321/
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture comprises are particularly locko promises are acquered to 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, ceramic agents and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface conducting used (1) for analysis of patterns, especially minerals, trace elements, organic conductives or supplements, especially minerals, trace elements, organic conductives or supplements, especially minerals, trace elements, organic conductivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cailes famino, carboxylic or fatty acid or their derivatives, salts and minerals inaging of the complete pattern of all nucleic acid solid in a cell, and conduction of differences in the nucleic acid properties o
                                                                                                                                                              useful for analyzing expression uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                capture probes used in the method of the invention.
                                                                                                                                                           Sorting single-stranded nucleic acid, patterns and screening active agents, and constant regions.
     (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                            Example; Page 5; 8pp; German.
                                                           Boekenkamp D, Dieck HT,
                                                                                                              WPI; 2003-714082/68.
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ô Gaps .; 0 Query Match 0.7%; Score 18; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 9.3e+02; Matches 18; Conservative 0; Mismatches 0; Indels Seguence 21 BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;

ADK01321 standard; DNA; 21 BP 06-MAY-2004 (first entry)

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

04-SEP-2003

Dieck HT, Hoppe H; (DEGS ) DEGUSSA BIOACTIVES GMBH. 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794 WPI; 2003-714082/68. Boekenkamp D,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded cucleic acids by isolation and hybridisation of nucleic acids pools, then reading out, where the nucleic acids are selectively bound using capture reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprises variable and non-variable regions. The capture oligonucleotides have a 5-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3-variable, discriminatory region that comprises a sequence of solid solid stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues The capture of capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, especial, and additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and elect very small differences in pools of nucleic acids from cells and comparison of allows inaspensal and reproducible. Since the can detect very small differences in the method spools, not individual center probes used in the method of the invention. This invention describes a novel method for sorting single-stranded

Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;

Gaps ; 0.7%; Score 18; DB 1; Length 21; 100.0%; Pred. No. 9.3e+02; 0; Indels 100.0%; Pred. ... Best Local Similarity 100.0 Matches 18; Conservative Query Match

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멾 ADK01322 standard; DNA; 21 (first entry) 06-MAY-2004 ADK01322; RESULT 1174 

Rat DNA microarray capture oligonucleotide #42.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

gg Rattus

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(DEGS ) DEGUSSA BIOACTIVES GMBH.
             28-FEB-2002; 2002DE-01008794.
                    28-FEB-2002; 2002DE-01008794.
                                                              Example; Page 5; 8pp; German.
                                   Boekenkamp D, Dieck HT,
                                                       and constant regions.
                                         WPI; 2003-714082/68.
DE10208794-A1.
       04-SEP-2003.
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Hoppe H;

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

This invertion describes a movet method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agence that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at certain act nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) conducting propertied by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, conducting broot, blood, nerve or germ cells and (ii) for date matrix is of for a crivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic crids and analysis of patterns, state and reproducible.

Conducting of the complete pattern of all nucleic acids from cells and callows imaging of the complete pattern of all nucleic acids from cell, and can lows imaging of the complete pattern of all nucleic acid pool: Since the method is based on comparison of nucleic acid pool: Since the can method is based on subplements appears the method is possible. The method is possible and the propertion of patterns and pools of the analysis of the analy genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent capture probes used in the method of the invention. This invention describes a novel method for sorting single-stranded

Sequence 21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;

0.7%; Score 18; DB 1; Length 21; 100.0%; Pred. No. 9.3e+02; 0; Indels 100.0%; Pred. ... Query Match
Best Local Similarity 100.
Matches 18; Conservative

2709 AAAAAAAAAAAAAA 2726 

ò g

ADK01324 standard; DNA; 21 BP. ADK01324; RESULT 1175 XXEXEXEX XXEXEXEX

Rat DNA microarray capture oligonucleotide #44.

06-MAY-2004 (first entry)

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;

blood; nerve; germ cell; food additive; food supplement.

Rattus sp.

DE10208794-A1.

04-SEP-2003.

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS ) DEGUSSA BIOACTIVES GMBH.

Hoppe H; Boekenkamp D, Dieck HT,

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions. WPI; 2003-714082/68.

Example; Page 5; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nuclectides to allow binding of particulars sorts of single stranded nucleic acids. The capture comprises as sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonuclectides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, physical, stimulated by an electrical field or through a molecular, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic crimistroot, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic caids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible

Compound of differences in pools of nucleic acid pool, since the method is based on comparison of nucleic acid pools, not individual canning and in the method is the physical in the nucl capture probes used in the method of the invention. 

Sequence 21 BP; 0 A; 2 C; 0 G; 19 T; 0 U; 0 Other;

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Gaps

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0.7%; Score 18; DB 1; Length 21; 00.0%; Pred. No. 9.3e+02; 0; Indels 100.0%; Pred. No. 7... rive 0; Mismatches 18; Conservative Best Local Similarity Query Match Matches

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Gaps ;

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ABD25933 standard; DNA; 21 ABD25933; RESULT 1176 ABD25933 SXXXE

29-JUL-2004 (first entry)

(EPIG-) EPIGENESIS PHARM INC WO200285309-A2 Homo sapiens. 31-OCT-2002 Miller S, Nyce JW, 

AA505075-derived oligonucleotide SEQ ID 4945.

beta-adrenergic agonist, respiratory disease; pulmonary vasoconstriction, respiratory distress syndrome; allergic rhinitis; pulmonary hypertension, emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human, antisense, bronchoconstriction, allergy; hyposecretion; pain, respiratory tract inflammation; adenosine sensitivity, lung, cancer; surfactant depletion; antiallergic, antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

Aguilar D; Pabalan J, Katz E, Sandrasagra A, Ka L, Shahabuddin S; Li Y, Sar Tang L,

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4945; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconscriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) receptors, confidenceonscriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) receptors, coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung evice, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antialfalmmatory, antiaethmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies as thma, impeded respiration, respiratory distress syndrome, path, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchit the breakced, the reduced adenosine content of the anti-sense oligos corresponding to the reduced adenosine content of the anti-sense oligos-corresponding to the oligonucleotides into products that fare adenosine into the system engine or in the target that fare adenosine into the system engine or in the target that fare adenosine or the edgradation and products that fare adenosine content of the edgradation benchman and the

Sequence 21 BP; 17 A; 0 C; 0 G; 4 T; 0 U; 0 Other;

Gaps ; 0 0.7%; Score 18; DB 1; Length 21; 00.0%; Pred. No. 9.3e+02; 0; Indels Best Local Similarity 100.0%; Pred. No. 9.3 Matches 18; Conservative 0; Mismatches Query Match

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Gaps

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0; Indels

Score 18; DB 1; Length 21; Pred. No. 9.3e+02;

Best Local Similarity 100.0%; Pred. No. 9.3 Matches 18; Conservative 0; Mismatches

Query Match

2709 AAAAAAAAAAAAAA 2726

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The invention relates to a class of CpG immunostimulatory
oligonuclectides containing a 5'TCG motif or a CG at or the 5' end that
coligonuclectides containing an immune response. Oligonuclectides and
compositions of the invention are useful for treating allergy or asthma,
viral and bacterial infections and cancer e.g. biliary tract cancer,
breast cancer, cervical cancer, choriocarcinoma, colon cancer,
endometrial cancer, gastric cancer, lymphomas, intraepithelial melanoma,
liver cancer, lung cancer (e.g. small cell and non-small cell), melanoma,
curoblastomas, ovarian cancer, pancreatic cancer, brostate cancer,
rectal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer,
cetal cancer, sarcomas, thyroid cancer, renal cancer, brain
and CNS cancer, connective tissue cancer, oseophageal cancer, brain
Hodgkin's lymphoma, larynx cancer, oral cavity cancer, skin cancer,
ceticular cancer, as well as other carcinomas and sarcomas. The
cincention is also useful in gene therapy. The present sequence is a CpG
immunostimulatory oligonucleotide.
                                                                                                                                                                                                                                                                            CpG immunostimulatory oligonucleotide; immune response; allergy; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 viral and
breast
                                                                                                                                                                                                                                                                                              viral infection; bacterial infection; cancer; lymphoma;
intraepithelial neoplasm; melanoma; neuroblastoma; Hodgkin's lymphoma;
carcinoma; sarcoma; gene therapy; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New oligonucleotides, useful for treating allergy or asthma, bacterial infections, and cancer, e.g. biliary tract cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Krieg AM, Jurk M, Vollmer J, Uhlmann E;
                                                                                                                                                                                                                                        CpG immunostimulatory oligonucleotide #13.
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                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/mod_base= OTHER
2708 TAAAAAAAAAAAAAA 2725
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25-SEP-2003; 2003US-0506108P.
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                                    4 TAAAAAAAAAAAAAAA
                                                                                                                            ADP86142 standard; DNA; 21
                                                                                                                                                                                                   09-SEP-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bacterial infections, and cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
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misc feature
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17-SEP-1993;
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                                                                                                                                                       28-APR-1994,
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         Synthetic.
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                               Key
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                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an ICAM-1 binding protein, and a polynucleotide encoding the same, a vector and a host cell containing the same polynucleotide, and a composition comprising the same protein are provided. The ICAM-1 binding protein has high specificity to ICAM-1 and small molecular size, so that it can be effectively expressed in Bacherichia coli. and the composition containing the ICAM-1 binding protein has improved stability and excellent tissue penetration, so that it can be useful for the treatment of inflammation. The present sequence represents a mouse ICAM-1 binding protein associated PCR primer.
                                                                                                                                                                                                                                                                                                                                          ICAM-1 binding protein, and polynucleotide encoding the same, vector and host cell containing the same polynucleotide, and composition comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2',5'-linked tetraadenylate-antisense oligonucleotide chimeric mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense; 2',5'-tetraadenylate; 2-5A dependent RNase activator; RNA cleavage; antiviral therapy; chimeric molecule; ss.
                                                                                                                    Mouse ICAM-1 binding protein associated PCR primer SEQ ID NO 10.
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                                                                                                                                         ICAM-1 binding protein; ICAM-1; inflammation; 88; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.7%; Score 18; DB 1; Length 21;
100.0%; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ilarity 100.0%; Pred. No. 9.3
Conservative 0; Mismatches
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                                                          BP
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                                                                                                                                                                                                                                                35-MAR-2003; 2003KR-00013610
       21 AAAAAAAAAAAAAA
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                                                         ADX69527 standard; DNA; 21
                                                                                                  21-APR-2005 (first entry)
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Matches 18; Conserv
                                                                                                                                                                                                                                                                    (HAHN/) HAHN J H.
                                                                                                                                                                                                                                                                                                    Lee WJ;
                                                                                                                                                                                                                                                                                                                                                               the same protein.
                                                                                                                                                                                                                                                                               (LEEW/) LEE W J.
                                                                                                                                                                                  KR2004078763-A.
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04-JAN-1995
                                                                                                                                                               Synthetic.
                                                                             ADX69527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                     RESULT 1178
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This sequence is an example of a 2-5A-antisense oligonucleotide chimeric molecule. The antisense region targets the chimeric molecule to a particular region of RNA to be specifically cleaved and the 2',5'-linked tetraadenylate tail activates the 2-5A RNase. Typical applications are treatment of viral infections (esp. for cleavage of an RNA virus genome), cancer, leukaemia, cardiovascular disorders (e.g. restenosis after angioplasty), genetic disorders, osteoarthritis or rheumatoid arthritis. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       opecific cleavage of RNA, useful partic. for treating viral infection, cancers, etc. - by using anti-sense oligo:nucleotide coupled to activator of 2-5A dependent RNase.
                                                                                                                                       phosphodiester bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                            /*tag= a //s.-linked tetraadenylate /label= 2,5'-linked tetraadenylate /note= "nucleotides linked through phosmat hydroxyl groups of 2' and 5' carbons'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 4 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                        /note= "antisense region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (USSH ) US DEPT HEALTH & HUMAN SERVICES. (CLEV-) CLEVELAND CLINIC RES INST.
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Location/Qualifiers
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93US-00123449
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JP06303997-A.
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                                            Query Match
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                                                                                                      Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded CDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of CDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
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                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                             y Match 0.6%; Score 17.8; DB 1; Length 21; Local Similarity 90.5%; Pred. No. 9.6e+02; hes 19; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                         Sequence 21 BP; 1 A; 0 C; 3 G; 17 T; 0 U; 0 Other;
                                                           (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                               2707 CTAAAAAAAAAAAAAAAA 2727
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                                                                                                                                           Disclosure; Page 5; 11pp; Japanese.
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             16-APR-1993;
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Matches
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                            Gaps
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method can be used to analyse gene expression rapidly and easily
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                                                                                                          0.6%; Score 17.8; DB 1; Length 2
90.5%; Pred. No. 9.6e+02;
/ative 0; Mismatches 2; Indels
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                                                     Sequence 21 BP; 2 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
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Pred. No. 9.6e+02;
0; Mismatches 2;
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nes 19; Conservative
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                                                                                                                                                                                                                                             double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The method can be used to analyse gene expression rapidly and easily
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                    Analysis; gene expression; reverse transcription; primer; cDNA;
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restriction enzyme; ss.
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Reverse transcription primer used in cDNA analysis technique.
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                             aggregate; restriction enzyme; ss
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                                                                               double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 9.6e+02;
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Disclosure; Page 8; 11pp; Japanese.
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gene expression - by amplification of mRNA followed
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                             Score 17.8; DB 1; Length 21;
Pred. No. 9.6e+02;
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GBNESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
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                                   A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Disclosure; Page 8; 11pp; Japanese.
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of ababiled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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0.6%; Score 17.8; DB 1; Length 21; 90.5%; Pred. No. 9.6e+02;
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                                                                 0; Mismatches
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aggregate; restriction enzyme; ss.
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           Length 21;
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         Query Match 0.6%; Score 17.8; DB 1; Best Local Similarity 90.5%; Pred. No. 9.6e+02; Matches 19; Conservative 0; Mismatches 2;
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Analysis of cDNA and gene expression - by amplification of mRNA followed
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-CD5798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and, (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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llarity 90.5%; Pred. No. 9.6e+02;
Conservative 0; Mismatches 2; Indels
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Pred. No. 9.6e+02;
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Synthetic.

AAQ75659;

RESULT 12 AAQ75659/

Query Match

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Matches

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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Reverse transcription primer used in cDNA analysis technique.
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Score 17.8; DB 1;
Pred. No. 9.6e+02;
); Mismatches 2;
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Pred. No. 9.6e+02;
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                    (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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Pred. No. 9.6e+02;
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                         - by amplification of mRNA followed
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0; Mismatches 2;
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labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                     Query Match
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                           reverse transcription; primer; cDNA;
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                                    Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         restriction enzyme; ss.
                                                                             yene expression; reverse restriction enzyme; ss
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                                                                           Analysis;
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AAQ75761 standard; DNA; 21

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ7547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                alysis of cDNA and gene expression - by amplification of mRNA followed digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression - by amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                0.6%; Score 17.8; DB 1; Length 21; 90.5%; Pred. No. 9.6e+02; ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           by digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                                                                                         2706 ACTAAAAAAAAAAAAAAA 2726
                                                                                                       Disclosure; Page 9; 11pp; Japanese.
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AAQ75654 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis;
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                                                                                                                                                                                                                                                               restriction enzyme; ss.
                                                      GGACAAAAAAAAAAAAA 1
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nes 19; Conserv
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19;
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This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is precancerous condition, by administering to the patient a first allale specific inhibitor (ASI) targeted to an allele of a first essential gene specific inhibitor (ASI) targeted to an allele of a first essential gene present in cells of the precancerous condition, where the normal somatic calls of the patient are heterozygous for the first gene, the inhibitor is active on at least one but less than all allelic forms present in the present in a population and targets only one allelic form present in the used in the diagnosis, prevention and treatment of LOH disorders, e.g. cancers, atherosclerotic plaques, premalignant metaplastic or dysplastic cancers, benign tumours, endometriosis, polycyclic kidney disease, and graft versus host disease. The method can also be used to remove mallymant cells from bone marrow transplants. AZ25812-Z26825 represent thus an polymorphic sites described in the method of the invention
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                                                                                                                                                                                                                                                                                                                                                 Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
dysplastic lesion, benign tumour, polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
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                                                                                                                                                                                                                                                                             Stanton VP;
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                                                                                                                                                                                                                                            (VARI-) VARIAGENICS INC.
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Matches 19; Conserv
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                                                           Homo sapiens
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The present invention relates to a novel pharmaceutical composition comprising: (a) an agent capable of modulating an expression level or protein activity of a cancer-related transmembrane protein (CRTP) or gene; protein activity of a cancer-related transmembrane protein (CRTP) or gene; an antibody specific for a CRTP, or a carrier. The pharmaceutical composition may also comprise a polymucleotide capable of inhibiting or decreasing the expression of the cryp paramaceutical composition may also comprise a polymucleotide capable of inhibiting or decreasing the expression of the invention are selected from ABCC4, C20orf103, CACNALD, CDH6, CST, ENPP3, FLJ1856, GPR54, HAVCR1, SLC6A3, SLC30A4, TRG, and TRPM4. The cancer, lung cancer, breast cancer, prostate cancer, lung cancer, breast cancer, prostate cancer, lung cancer, breast cancer, prostate cancer, liver cancer, stomach cancer, and esophageal cancer. The present sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at tp. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                    Pharmaceutical composition comprising an agent capable of modulating an expression level or protein activity of a gene, e.g. ABCC4, or a T cell activated by the polypeptide or antibody, and a carrier, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                              Claim 3; SEQ ID NO 14541; 113pp; English.
                                                                                                                     SH;
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              19-MAY-2004; 2004WO-US015645.
                                               20-MAY-2003; 2003US-0471729P
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                                                                                                                   Wei L, Slonim DK,
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                                                                                                                                                                                                                                               treating cancer.
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                                                                                    (AMHP ) WYETH
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The present sequence is a telomerase Oliog-dT-Primer, which can be used in a novel method for detecting telomerase activity. The method comprises adding to a test sample a 1st primer, that serves as telomerase adding to a test sample a 1st primer, that serves as telomerase substrate, and nucleoside triphosphate (dNTP) and incubating to allow primer extension by the telomerase, amplifying the extension product, immobilising the amplification product (AP) on a solid phase and qualitative and/or quantitative detection of AP, where the substrate primer is preferably from the 5-region of the long terminal repeat 2 (LTR-2) sequence of a retrovirus. The method can be used to diagnose tumours and screen compounds for effector activity. Immobilisation of AP provides a signal that is reproducibly representative of telomerase activity, eliminates the need for gel electrophoretic separation and provides high sensitivity. Radioactive labels are not required and the method can be automated for routine uses. Specific detection is achieved by proper choice of hybridisation conditions, without separation of the telomerase extension product. A specific signal is generated by 1-10 cell equivalents, but for tumour analysis 10-1000 ng of tissue is usually used
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                                                           Measuring telomerase activity, useful for tumour diagnosis and compound screening - by extending substrate primer, followed by amplification and immobilising product for detection.
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nes 17; Conservative
                     WPI; 1997-299542/28.
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and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                         Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; 88.
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                                      Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 17.4; DB 1;
94.7%; Pred. No. 9.7e+02;
iive 0; Mismatches 1;
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                                         digestion with restriction enzymes.
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                                                                                                          Disclosure; Page 5; 11pp; Japanese.
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          of cDNA and
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The invention relates to detecting mutations in the BRCA1 and hMLH1 gene comprising subjecting a set of amplification products to two-dimensional DNA electrophoresis (TGDS) to produce a characteristic spot pattern for a specific mutation in either the BRCA1 or the hMLH1 gene. Also included are test kits for enabling BRCA1 or hMLH1 gene testing comprising short KCD, 25 micro M of dNTP, and 5 % formamide. The method is useful for detecting mutations in the BRCA1 (breast and ovarian cancer susceptibility gene, a tumour suppressor gene) and hMLH1 gene (a DNA mismatch repair gene). The present sequence is a PCR primer specific to hMLH1 used in the method of the invention
                                                                                                                                                                                                                                                                                                               Detecting mutations in the BRCA1 and hMLH1 gene comprises subjecting amplification products to 2-dimensional gel electrophoresis to produce characteristic spot pattern for a specific mutation in either the BRCA1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression; reverse transcription; primer; cDNA;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                        or the hMLH1 gene
                                      WO200236819-A1.
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  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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              2707 CTAAAAAAAAAAAAAA 2725
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- by amplification of mRNA followed
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                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded CDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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0.6%; Score 17.4; DB 1; Length 20;
Best Local Similarity 94.7%; Pred. No. 9.9e+02;
Matches 18; Conservative 0; Mismatches 1; Indels
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0.6%; Score 17.4; DB 1; Length 20;
Best Local Similarity 94.7%; Pred. No. 9.9e+02;
Matches 18; Conservative 0; Mismatches 1; Indels
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  Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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                                                                Disclosure; Page 5; 11pp; Japanese.
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Pred. No. 9.9e+02;
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Best Local Similarity
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Pred. No. 9.9e+02;
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                                                                                                                                                                                                                                                                                                           double-stranded cDNAs by using an aggregate of mRNAs and a pluzil type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                       (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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Best Local Similarity 94.7%;
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                  Analysis, gene expression, reverse transcription; primer, cDNA; aggregate, restriction enzyme; ss.
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Reverse transcription primer used in cDNA analysis technique.
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Pred. No. 9.9e+02;
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Best Local Similarity 94.7
Matches 18; Conservative
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ilarity 94.7%; Pred. No. 9.9e+02;
Conservative 0; Mismatches 1; Indels
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Local Similarity 94.7%; Pred. No. 9.9e+02;
les 18; Conservative 0; Mismatches 1; Indels
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                                                                      double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GBNESGO files AAQYIS47-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 9.9e+02;
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                   Page 5; 11pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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Best Local Similarity
Matches 18; Conserva
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                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1250
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                                                                                                                                                                                                                                                                                                    Matches
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The present sequence is an oligomucleotide provided in a specification relating to the simplified purification of oligonucleotides by liquid chromatography. The process comprises: (a) pouring oligonucleotides of chromatography. The process comprises: (a) pouring oligonucleotides of group into a liquid chromatography column packed with an acid and alkaling resistant packing agent, such as polystyrene resin; (b) pouring a mixed developing solvent composed of a buffer made from a volatile salt and a contract soluble organic solvent at a suitable concentration gradient into the column; (c) pouring an acid, particularly 6-16 v/v% acetic acid, into the column to deprotect the oligonucleotides protected with the chydrophobic group; (d) pouring a mixed developing solvent composed of a buffer made from a volatile salt, particularly 0.05-0.5 N aqueous camponium hydrogencarbonate solution adjusted at pH 8-10, and a water soluble organic solvent at a suitable concentration gradient to elute the deprotected oligonucleotides; and (e) removal of the solvent and the salt from the eluted oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; cancer; autoimmune disease; organ transplantation; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human dendritic cell wall membrane molecule-associated primer #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.6%; Score 17.4; DB 1; Length 20;
14.7%; Pred. No. 9.9e+02;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           process for purification of oligonucleotides using liquid
                                                                                                                 Oligonucleotide purification; liquid chromatography; hydrophobic protective group; deprotection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 17 A; 1 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 4; 13pp; Japanese.
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                                                                                   Synthetic oligonucleotide #9.
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                                                   entry)
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Best Local Similarity 94.7%
                                                 (first
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                                                                                                                                                                                                    JP2000342265-A.
                                                                                                                                                                                                                                                                                                                                                                                                                             chromatography.
                                                                                                                                                                                                                                                                       02-JUN-1999;
                                                                                                                                                                                                                                                                                                        02-JUN-1999;
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                                                 12-JUL-2001
                                                                                                                                                                                                                                     12-DEC-2000.
                                                                                                                                                                    Synthetic.
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                                                                                                                                                                                                                                                                                              The invention relates to an isolated human dendritic cell wall membrane molecule comprising a defined amino acid sequence given in the specification, or its variant based on the amino acid sequence but with some amino acids deleted, substituted, inserted and/or added and capable of controlling immune response. The protein, variants and encoded DNNs are useful in producing antibodies and soluble molecules to separate or detect dendritic cells, and for treatment of cancer, autoimmune diseases, organ transplantation, infection and allergy, e.g. by cancer vaccines and dendritic cell therapy to control immune response through promotion or buman dendritic cell wall membrane increases expression with maturation of human dendritic cells and membrane sequence represents a human dendritic cell wall membrane molecule-associated primer
                                                                                                                                                                                                                    Isolated dendritic cell wall membrane, variants and their encoded DNAs, useful in producing antibodies and soluble molecules to separate or detect dendritic cells, and for treatment of cancer, autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; cancer; autoimmune disease; organ transplantation; infection; allergy; vaccine; dendritic cell therapy; immune response; primer; ss; dendritic cell wall membrane molecule; immunogenic.
allergy; vaccine; dendritic cell therapy; immune response; primer; s8; dendritic cell wall membrane molecule; immunogenic.
                                                                                                                                                                         Imai N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human dendritic cell wall membrane molecule-associated primer #1.
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                                                                                                                                                                         Ehara H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.6%; Score 17.4; DB 1; Length 20;
14.7%; Pred. No. 9.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                         Hinohara A, Nakagawa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 3 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                               Example 6; Page 20; 68pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2094 ACCCCGTGCTGCAGCAGCA 2112
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                                                                                                      12-SEP-2001; 2001WO-JP007919.
                                                                                                                             12-SEP-2000; 2000JP-00277352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ilarity 94.7%;
Conservative
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                                                                                                                                                                         Watarai H, Yamaguchi Y,
                                                                                                                                                  (KIRI ) KIRIN BEER KK
                                                                                                                                                                                                WPI; 2002-362337/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 18; Conserv
                                                          WO200222683-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200222683-A1
                                                                                                                                                                                                                                                         and infection.
                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                 21-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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ABK48093
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      molecule comprising a defined amino acid sequence given in the specification, or its variant based on the amino acid sequence but with some amino acids deleted, substituted, innerted and/or added and capable of controlling immune response. The protein, variants and encoded DNAs are useful in producing antibodies and soluble molecules to separate or detect dendritic cells, and for treatment of cancer, autorimune diseases, organ transplantation, infection and allergy, e.g. by cancer vaccines and suppression of the interaction between dendritic cells and T cells. The human dendritic cell wall membrane increases expression with maturation of human dendritic cells. The present sequence represents a human dendritic cells. The sequence represents a human dendritic cell wall membrane molecule-associated primer
                                                                                                                                                                                                                                              Isolated dendritic cell wall membrane, variants and their encoded DNAs, useful in producing antibodies and soluble molecules to separate or detect dendritic cells, and for treatment of cancer, autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, antisense, lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to an isolated human dendritic cell wall membrane
                                                                                                                        Imai N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                            Ehara H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
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                                                                                                                            Nakagawa R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4; DB 1;
Pred. No. 9.9e+02;
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                                                                                                                            Hinohara A,
                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 20; 68pp; Japanese.
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Tang L, Shahabuddin S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.6%;
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12-SEP-2000; 2000JP-00277352
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                                                                                                                            χ,
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                                                                                                                            Yamaguchi
                                                                  (KIRI ) KIRIN BEER KK
                                                                                                                                                                                        WPI; 2002-362337/39.
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                                                                                                                                                                                                                                                                                                                                                 and infection.
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                                                                                                                               Watarai H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Miller S,
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Pharmaceutical composition for treating allments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 776; 872pp; English.

first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antisinflammatory steroid and ubiquinone. A composition of the invention chas antisinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or mallignant crespiratory effect of an entiallammatory steroid in a subject, for reducing levels of for enhancing the prophylactic or therapeutic respiratory effect of an entiallammatory steroid in a subject, for reducing levels of denosine creceptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung suffactant in a subject is tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the content of the printed and the prin The invention relates to a novel pharmaceutical composition, which has at ftp.wipo.int/pub/published\_pct\_sequences

20 BP; 18 A; 0 C; 2 G; 0 T; 0 U; 0 Other; Sequence

Gaps .. 0 0.6%; Score 17.4; DB 1; Length 20; 94.7%; Pred. No. 9.9e+02; ive 0; Mismatches 1; Indels 1; Indels 2709 AAAAAAAAAAAAAAA 2727 Local Similarity 94.7 nes 18; Conservative Query Match Best Local S Matches ò

AAAAAAAAAAAAAA 19

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Human oligonucleotide sequence ABZ89487 standard; DNA; 20 BP (first entry) 17-OCT-2003 ABZ89487; RESULT 1255 ABZ89487 

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Œ, Katz ii Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Miller S, ₹

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 4729; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antisinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing bronchodilation, increasing levels of ubiquinone or lung suffactant in a subject's tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Under all maniformation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cut fitp.wipo.int/pub/published\_pct\_sequences

Sequence 20 BP; 18 A; 2 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Score 17.4; DB 1; Length 20; Pred. No. 9.9e+02; 0; Mismatches 1; Indels 0; Mismatches 0.6%; 18; Conservative Local Similarity Query Match Matches

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ABZ92865 standard; DNA; 20 (first entry) 17-0CT-2003 ABZ92865; RESULT 1256 

ВР.

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Katz ŝ Li Y, Sandrasagra A, Tang L, Shahabuddin Nyce JW, I Miller S,

Pabalan J, Aguilar D;

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WPI; 2003-229219/22.

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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                ubiquinone
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Disclosure; SEQ ID NO 8107; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intistion codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, 5' or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or casal alraway dysfunction and a second active agent comprising an antinflammatory steroid and ubiquinone. A composition of the invention has antinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also antinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of an antilatematory broncholiation, increasing levels of adenosine or lung surfactant in a subject, s tissue, or treating bronchoconstriction, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed condition. specification, but was obtained in electronic format directly from WIPO ftp.wipo.int/pub/published\_pct\_sequences 

Sequence 20 BP; 18 A; 0 C; 1 G; 1 T; 0 U; 0 Other;

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                                Gaps
                                .;
0.6%; Score 17.4; DB 1; Length 20; 94.7%; Pred. No. 9.9e+02;
                             1, Indels
                              0; Mismatches
 Query Match
Best Local Similarity 94.7
Matches 18; Conservative
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2706 ACTAAAAAAAAAAAAAA 2724 AGTAAAAAAAAAAAAA 20 ઠે 셤

ABZ88938 standard; DNA; 20 BP (first entry) 17-0CT-2003 ABZ88938; RESULT 1257 ABZ88938 

Human oligonucleotide sequence.

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antisethmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC

Aguilar D; Pabalan J, Katz E, Sandraвagra A, Ka L, Shahabuddin S; Li Y, Sar Tang L, Miller S, Nyce JW,

WPI; 2003-229219/22

WPI; 2003-229219/22

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a condition, as unbined in a subject, for reducing or depleting levels of for entitinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing levels of adenosine cuse in a subject, a tissue, or treating bronchoconstruction, cuse in a time and part in a subject is tissue, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed process or the condition of the cond ö Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or Gaps ö 0.6%; Score 17.4; DB 1; Length 20; 4.7%; Pred. No. 9.9e+02; 1; Indels Sequence 20 BP; 17 A; 1 C; 1 G; 1 T; 0 U; 0 Other; at ftp.wipo.int/pub/published\_pct\_sequences Disclosure, SEQ ID NO 4180; 872pp; English. 0; Mismatches 2708 TAAAAAAAAAAAAAA 2726 2 TCAAAAAAAAAAAAAA 20 Human oligonucleotide sequence. BP. 94.78; ABZ89872 standard; DNA; 20 (first entry) Best Local Similarity 94.7 Matches 18; Conservative 17-0CT-2003 ubiquinone ABZ89872; Query Match RESULT 1258 ABZ89872 요 8

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiinsthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. Aguilar D; Katz E, Pabalan J, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013135. 24-APR-2001; 2001US-0286137P. (EPIG-) EPIGENESIS PHARM INC WO200285308-A2. Homo sapiens 31-OCT-2002. Miller S, Nyce JW, 

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Disclosure; SEQ ID NO 5114; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, 5' or zegions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or masal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, cus in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or therapeutic respiratory effect of an autisinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine treeptor, producing bronchodilation, increasing levels of ubiquinone or lung suffammation, lung allergies, or a respiratory disease or condition, lung autisation, and a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fire, wipo int/pub/published\_pct\_sequences

Sequence 20 BP; 16 A; 2 C; 1 G; 1 T; 0 U; 0 Other;

Gaps . 0 DB 1; Length 20; 1; Indels Score 17.4; DB 1; Pred. No. 9.9e+02; 0; Mismatches 0.6%; 18; Conservative Local Similarity Query Match Matches

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ò 요 RESULT 1259 ABD26102

ABD26102 standard; DNA; 20 BP ABD26102; 

29-JUL-2004 (first entry)

AA463249-derived oligonucleotide SEQ ID 5114.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

sapiens. Homo WO200285309-A2.

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC

Pabalan J, Aguilar D; Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Nyce JW, I Miller S,

WPI; 2003-093058/08.

oligonucleotide containing less percentage of adenosine, targeted t nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. composition for treating asthma, has antisense Pharmaceutical

Claim 15; SEQ ID NO 5114; 763pp; English.

The invention describes a novel composition, all eviating decrive agence, comprising oligomucleotides, effective for alleviating decribes and describes, effective for alleviating comprising oligomucleotides, effective for definition, allergies and bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligomucleotides are derived from a gene encoding or regulating creating a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligomucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analyses or composition or cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The culture of inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, willergies and/or surfactant hypoproduction are associated whypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cost criminamation, allergies and the manual proportion are associated transplantation rejection, pulmonary impeded respiratory.

Chypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cost transplantation rejection, pulmonary infections by the prevent the breakdown of the oligonucleotides into products that free addenosine content of the anti-sense oligonucleotides into products that fre e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it This invention describes a novel composition (a) a first active agent,

Seguence 20 BP; 16 A; 2 C; 1 G; 1 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 17.4; DB 1; Length 20; 94.7%; Pred. No. 9.9e+02; ive 0; Mismatches 1; Indels 1 Similarity 94.7 Query Match Local Matches

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2707 CTAAAAAAAAAAAAAA 2725 CTCAAAAAAAAAAAAAA 19 Н

ð 셤 RESULT 1260 ABD21764

ABD21764 standard; DNA; 20 BP

ABD21764; 

(first entry) 29-JUL-2004 Human stanniocalcin-derived oligo SEQ ID 776.

analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allorgic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation; adenosine sensitivity; lung; cancer, surfactant depletion; antiallergic; antiinflammatory; antiasthmatic;

sapiens Homo ngs.res

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoonstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configuration or desirved from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung cartering also describes a kit, that comprises: (a) a delivery device, in invention also describes a kit, that comprises: (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiaflammatory, antiathmaticy, is a nalgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung and is allergic finilammation, allergies and/or bronchoconstriction and as associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory chierress syndrome, pain, cystic fibrosis, allergic thinits, pulmonary chierress syndrome, pain, cystic fibrosis, allergic thinits, pulmonary chierrension, emphysema, chronic observation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
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                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted inucleic acids associated with lung airway or lung dysfunction, and
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Shahabuddin
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Best Local Similarity 94.7%;
Marches 18; Conservative
                                                                                                 23-APR-2002; 2002WO-US013143
                                                                                                                                               24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                 Sandrasagra
                                                                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC
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Tang L,
WO200285309-A2.
                                                31-OCT-2002
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ö Gaps ö 0.6%; Score 17.4; DB 1; Length 20; 4.7%; Pred. No. 9.9e+02; Indels

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AI034360-derived oligonuclectide SEQ ID 4729 ABD25717 standard; DNA; 20 (first entry) 29-JUL-2004 ABD25717; RESULT 1261 2×2×5×3

respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cytic fibrosis; beta-adrentatic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Human; antisense; bronchoconstriction; allergy; hyposecretion; pain;

Homo sapiens. 

Aguilar D;

WO200285309-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

t 0

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Ğ, Nyce

WPI; 2003-093058/08.

t C Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 4729; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, of the invention has antiallergic, antiinflammatory, antiasthmatic, is a composition to a useful for preventing or treating a respiratory, lung or malignant disease. The administered composition are associated or composition composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition and/or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or surfactant hypoproduction and/or lung corresponding to treasplantation rejection, pulmonary vasoconstriction, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligon sorresponding to the main-sense oligon sense that free adenosine ento the system corrent of the anti-sense oligon sense and entous the target RNA serves to prevent the braskdown of the course of the oligonucleotides into products that free adenosine environment and thereby, the course of the oligonucleotides into product that the target RNA serves construction and prevent any unwanted effects due to it

Sequence 20 BP; 18 A; 2 C; 0 G; 0 T; 0 U; 0 Other;

Gaps ö Query Match 0.6%; Score 17.4; DB 1; Length 20; Best Local Similarity 94.7%; Pred. No. 9.9e+02; Matches 18; Conservative 0; Mismatches 1; Indels

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2706 ACTAAAAAAAAAAAAA 2724

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2 ACCAAAAAAAAAAAA 20

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RESULT 1262

ABD25168 standard; DNA; 20 BP

ABD25168;

29-JUL-2004 (first entry)

AI041482-derived oligonucleotide SEQ ID 4180.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

WO200285309-A2

31-OCT-2002

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

ä Aguilar Pabalan J, EI, Katz i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; ij Miller S, Š 

WPI; 2003-093058/08.

t C Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted truncleic acids associated with lung airway or lung dysfunction, and bronchodilating agent

Claim 15; SEQ ID NO 4180; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating by the comprising oligonucleotides, effective for alleviating by the constraint of pronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung capturction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention as antiallergic, antinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to composition, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory disease, pulmonary contains or an allergies, asthma, impeded respiration, respiratory continument of the anti-sense oligos corresponding to the ranget RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of

ö t 2 the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it Gaps ö Score 17.4; DB 1; Length 20; Pred. No. 9.9e+02; 1; Indels Sequence 20 BP; 17 A; 1 C; 1 G; 1 T; 0 U; 0 Other; 0; Mismatches Pred. No. 0.6%; 94.78; 18; Conservative Similarity Query Match Local Matches SXSSS ò

2708 TAAAAAAAAAAAAAA 2726 TCAAAAAAAAAAAAAAA 20

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1263 ABD2909

BP ABD29095 standard; DNA; 20

ABD29095;

(first entry) 29-JUL-2004 AA679352-derived oligonucleotide SEQ ID 8107.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cycostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fihitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. 

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143..

24-APR-2001; 2001US-0286036P.

(EPIG-) EPIGENESIS PHARM INC.

Pabalan J, Aguilar D; Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Nyce JW,

WPI; 2003-093058/08

S Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 8107; 763pp; English.

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of the sensitivity and administered to a mammal. The surfactant depletion or hyposecretion, when administered to a mammal. The cligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a managesic. beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered

composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchocomstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, condition such as pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, oystic fibrosis, allergic rhintis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary characterion rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the milgonuclectides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it ö to nucleic acids encoding a mammalian glucocorticoid receptor. The antiense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone. New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome. antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE. The invention comprises an antisense oligonucleotides that are targeted Human glucocorticoid receptor-specific antisense oligonucleotide #3493. Gaps ; 0 0.6%; Score 17.4; DB 1; Length 20; 14.7%; Pred. No. 9.9e+02; Indels Sequence 20 BP; 18 A; 0 C; 1 G; 1 T; 0 U; 0 Other; 0; Mismatches Claim 4; SEQ ID NO 3493; 985pp; English 2706 ACTAAAAAAAAAAAAAA 2724 AGTAAAAAAAAAAAAA 20 ADH66659 standard; DNA; 20 BP. 20-MAY-2002; 2002US-0381857P. 20-MAY-2003; 2003WO-US016084 ilarity 94.7%; Conservative (first entry) Nalseth AE; (PHAA ) PHARMACIA CORP. WPI; 2004-035034/03 Best Local Similarity Matches 18; Conserv WO2003099215-A2. ното варіепв 25-MAR-2004 04-DEC-2003 Crosby SD, Query Match ADH66659; RESULT 1264 ADH66659, \$ a ð

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The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound such compound and composition are useful for treating a disease or condition compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, compound in trom burns, migraine headache, cluster headache, mild-to-moderate chadache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present compounds a chimeric phosphorothioate oligonucleotide with a deoxy gap. Used during the antisense inhibition of the human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                         Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; 88.
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             Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                          Score 17.4; DB 1;
Pred. No. 9.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberds SL;
                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                     ADK74413;
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                                            Query Match
                                                                                                                                                                                      RESULT 1265
                                                                Best Loca
Matches
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Matches
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microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1; inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; noctropic; antiathritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer; disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9g34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal antisense or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid
encoding mPGES-1, useful for preparing a composition for treating e.c
inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "phosphorothioate linkages and all cytidine residues are 5-methylcytidines"
                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:558
                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
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                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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/mod_base= OTHER
"^'-0-meth
                                                                                                                                                                                                                                                                                                                                                                                                                      mod_base= OTHER
   BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-2002; 2002US-0413549P.
ADM14371 standard; DNA; 20
                                                                    (first entry)
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*tag= b
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modified_base
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                                                                     01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                Synthetic
                                   ADM14371;
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ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diaberes, cancer, ischaemia or repertusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises antisense oligonucleotides that are targeted to the nucleic acids encoding a family of human proteins from mitochondrial membranes, which bind insulin sensitising, antidiabetic thiazolodinediones (referred to as: mitoNEET). The antisense oligonucleotides of the invention are useful for modulating mitoNEET expression and for treating diseases or conditions associated with mitoNEET, such as: diabetes, immunological disorders, cardiovascular disorders including hypertension, neurological disorders, and sichaemia/reperfusion injuries. The present DNA sequence represents a mitoNEET-specific antisense oligonucleotide of the invention. NOTE: The present sequence is a 2'-methoxyethyl (2'-MOE) gapmer with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                 0.6%; Score 17.4; DB 1; Length 20; 94.7%; Pred. No. 9.9e+02; tive 0; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; antisense oligonucleotide; mitochondrial membrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mitoNEET-specific antisense oligonucleotide #199.
                                                                                                                                           Sequence 20 BP; 0 A; 1 C; 2 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 2 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17.4; DB 1;
Pred. No. 9.9e+02;
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                                                                                                                                                                                                                                                              2707 CTAAAAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                      19 CCAAAAAAAAAAAAAA 1
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phosphorothioate backbone.
                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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RESULT 1268

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Matches

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)
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                                                                                                                                                                                                                                     Analysis of cDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                               A method for the analysis of cDNA comprises (a) preparing an aggregate c double-stranded cDNAs by using an aggregate of mRNAs and a plural type c labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) the double-stranded aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.4; DB 1
Pred. No. 1e+03;
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                                                                                                                                                                (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
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by digestion with restriction enzymes.
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Best Local Similarity 94.7
Matches 18; Conservative
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               JP06303997-A.
                                                                                        16-APR-1993;
                                                                                                                             16-APR-1993;
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                                                    01-NOV-1994.
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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                       expression; reverse transcription; primer; cDNA,
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                                                                                                                                                                                                                                                                                  Reverse transcription primer used in cDNA analysis technique.
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0.6%; Score 17.4; DB 1; Length 21;
Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismatches 1; Indels
Indels
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                           aggregate; restriction enzyme; ss
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                                     2708 TAAAAAAAAAAAAAAAA 2726
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                                                                                                                                                                  AAQ75622 standard; DNA; 21
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 Conservative
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   18;
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RESULT 1269

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Gaps

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RESULT 1271 AAQ75762/c

Matches

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AAQ75631;

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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A method for the analysis of cDNA comprises (a) preparing an aggregate
                                                                                                                   gene expression; reverse transcription; primer; cDNA;
restriction enzyme; ss.
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                                                                                  Reverse transcription primer used in cDNA analysis technique
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4.7%; Pred. No. 1e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 1 A; 0 C; 2 G; 18 T; 0 U; 0 Other;
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                                             (first entry)
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nes 18; Conservative
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                                                                                                                     Analysis; gene
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                                                                                                                                         aggregate;
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of
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and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                  Score 17.4; DB 1; Length 21;
Pred. No. 1e+03;
0; Mismatches 1; Indels
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94.7%; Pred. No. 1e+03;
Ive 0; Mismatches 1; Indels
                                                                                                                                                                                         1; Indels
                                                                                                                 Sequence 21 BP; 0 A; 2 C; 2 G; 17 T; 0 U; 0 Other;
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AAQ75762 standard; DNA; 21 BP.
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Best Local Similarity
Matches 18; Conserv
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                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                         Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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0.6%; Score 17.4; DB 1; Length 21;
Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismatches 1; Indels
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94.7%; Pred. No. 1e+03;
lve 0; Mismatches 1; Indels
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                                                                            Disclosure; Page 5; 11pp; Japanese
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Matches 18; Conservative
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WPI; 1995-018287/03
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                expression; reverse transcription; primer; cDNA;
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Pred. No. 1e+03;
0; Mismatches 1; Indels
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2708 TAAAAAAAAAAAAAAA 2726
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nes 18; Conservative
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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  transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                         Sequence 21 BP; 1 A; 2 C; 1 G; 17 T; 0 U; 0 Other;
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                                                                                                                                               0.6%; Score 17.4; DB 1 larity 94.7%; Pred. No. 1e+03; Conservative 0; Mismatches
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Best Local Similarity
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les 18; Conserv
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                                                                                                                                                      alysis of cDNA and gene expression - digestion with restriction enzymes.
                                                                                                                                                                            Disclosure; Page 8; 11pp; Japanese.
                                 gene expression; reverse; restriction enzyme; ss.
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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Mismatches 1; Indels
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Matches 18; Conserv
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Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                    Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                 Reverse transcription primer used in cDNA analysis technique.
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                   The
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 the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. method can be used to analyse gene expression rapidly and easily
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Pred. No. 1e+03;
0; Mismatches 1; Indels
                                                                                                               0.6%; Score 17.4; DB 1; Length 21; 94.7%; Pred. No. 1e+03; ative 0; Mismatches 1; Indels
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Best Local Similarity 94.77
Matches 18; Conservative
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Best Local Similarity 94.7
Matches 18; Conservative
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
 gene expression - by amplification of mRNA followed
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                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abablled reverse transcription primers (GENESSOR files AAQ75547.075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Pred. No. 1e+03;
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               digestion with restriction enzymes.
                                                     Disclosure; Page 6; 11pp; Japanese.
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AAQ75759 standard; DNA; 21 BP.
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Matches 18; Conservative
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Analysis of cDNA and gene expression - by amplification of mRNA followed
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Pred. No. 1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                               by digestion with restriction enzymes.
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                                                                                                                                                                                                    restriction enzyme; ss
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Best Local Similarity
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0.6%; Score 17.4; DB 1; Length 21; 94.7%; Pred. No. 1e+03; tive 0; Mismatches 1; Indels

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04-AUG-1995 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ75766;
                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1295
                                                                                                                                                                                                                                                                                     Matches
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This invention relates to novel genes and the encoded proteins thereof
that are derived from the sea squirt Ciona intestinalis. Specifically, it
refers to those genes that are expressed in the tissues or organs of the
sea squirt during its developmental phase. The present invention
c describes the identification of these genes as useful for elucidation of
the mechanism of development and hence for developing regeneration
c medicines and gene therapy techniques. Accordingly, they can be used in
the research of various genetic diseases, as well as the analysis of cell
proliferation, differentiation and reproduction. Furthermore, such
compositions can be useful for environmental measurements and water
c surveys, particularly for sea water surveys, and also for the preparation
of transformed sea squirt for improving edibility of sea squirt such as
thalocynthia roretzi. This oligonucleotide sequence is a PCR primer used
to amplify sea squirt DNA given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                        Novel gene cluster which is specifically expressed in tissue or organ during developmental phase of sea squirt, useful for elucidation of mechanism of development of tissue or organ of sea squirt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer; detection; characterisation; mRNA; restriction display PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 17.2; DB 1; Length 18; 94.4%; Pred. No. 9.7e+02; ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Template mRNA poly-A tail SEQ ID NO:1 from WO9729211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                             (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 38; 1846pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2708 TAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВЪ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-US002009.
                                                                                                                          31-JUL-2002; 2002JP-00222593
                                                                                                                                                                    31-JUL-2002; 2002JP-00222593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT94431 standard; mRNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthesis; cDNA; ss.
                                                                                                                                                                                                                                                     WPI; 2004-287079/27.
Ciona intestinalis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                           JP2004057129-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Weinstein JN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-FEB-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT94431;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents a primer used in the preparation of the DNA construct of the invention. The construct is for inserting a DNA fragment of interest into a eukaryotic host cell, and comprises an integration cassette flanked by site-specific recombination sequences in which is inserted the DNA of interest, where the DNA fragment of interest is flanked by a nucleotide sequence sharing homology to a nucleotide sequence sharing homology to a nucleotide sequence copy in the eukaryotic cell. The construct is used for genetic modification of diploid plant and animal cells. The integration cassette improves the genomic insertion of the DNA fragment of interest in a site-specific manner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA construct for genetic modification of eukaryotic cells - comprising integration cassette flanked by site-specific recombination sequences.
                                                                                                                                                                                                                                                                               PCR primer; integration cassette; site-specific recombination sequence; genetic modification; site-specific genomic insertion; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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cell proliferation; differentiation; reproduction;
environmental measurement; water survey; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 17.4; DB 1; Length 21; 94.7%; Pred. No. 1e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 8 C; 3 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer 1 used to amplify sea squirt DNA
                                                                                                                                                                                                                                        Primer KC-1 for vector construction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 17; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          275 ATTGAGGAATTTGGGGAGG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pothier F;
      19 TGAAAAAAAAAAAAAA 1
                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADP04929 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-CA000375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-00656838
                                                                                                         AAV17253 standard; DNA; 21
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                                                                                                                                                                                              (first entry)
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Best Local Similarity 94.77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sirard M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-042199/04
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                                                                                                                                                                                              28-MAY-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-1997
                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                  AAV17253;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gagne M,
                                                                   RESULT 1296
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Gaps

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Tue Nov

WPI; 1997-415362/38

Detection and characterisation of mRNA by restriction display PCR - comprising synthesis of cDNA, digestion with a restriction endonuclease, ligation to an adaptor DNA and PCR amplification.

Disclosure; Page 24; 40pp; English

A method has been improved for detecting and characterising mRNA colecules which includes synthesising a double stranded (ds) CDNA from molecules which includes synthesising a double stranded (ds) CDNA fromment comparison to an adaptor DNA sequence. The produce cDNA fragments in which at least one end of the CDNA fragments con end of the CDNA fragments. The major of the CDNA fragments to at least one end of the CDNA fragments; (b) ligating the adaptor DNA sequences to the CDNA fragments; (c) amplifying the cDNA fragments to at least one end of the CDNA fragments; (b) argainst that hybridise to the ends of the CDNA fragments, where the primers that hybridise to the ends of the CDNA fragments, where the primers have at least one nucleotide at the 3' end that specifically hybridises to a subset of CDNA molecules; and (d) detecting the presence of the resulting amplified CDNA fragments. The present sequence represent a template poly-A tail used in the present specification. The method designate restriction display PCR can be used for characterising cells based on their mNAA content, for representing expression. The method is also useful for characterising cells of a content of types and under a variety of physiological conditions. The cented of the content of is also useful for tissue from particular individuals or species based on the fingerprint obtained from the mRNA content of isolated cells or tissue and comparing it to cells or tissue

Sequence 19 BP; 17 A; 0 C; 0 G; 0 T; 0 U; 2 Other;

Gaps ö Score 17.2; DB 1; Length 19; 0; Indels 1e+03; Pred. No. 1e+0 1; Mismatches 0.6%; 94.48; Local Similarity 94.4 Query Match Matches

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AAX18390 standard; DNA; 19 BP. AAX18390, RESULT 1299 AAX18390, 

(first entry) 11-MAY-1999

RT-PCR primer of the invention SEQ ID 31.

RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.

Synthetic

JP11032765-A

09-FEB-1999

97JP-00208312 18-JUL-1997;

97JP-00208312 18-JUL-1997;

(TAKI ) TAKARA SHUZO CO

WPI; 1999-183822/16.

Peptides having at least two new nucleotides - useful as primers in RT-

Example 1; Page 12; 19pp; Japanese

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This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(apha)n-beta-N3'; or (X)m5'-(gamma) k-delta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thywine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine, cytosine or thymine; adenine; a natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
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Sequence 19 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 2 Other;

Gaps ö Query Match

0.6%; Score 17.2; DB 1; Length 19;
Best Local Similarity 94.4%; Pred. No. 1e+03;
Matches 17; Conservative 1; Mismatches 0; Indels

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RESULT 1300

AAA25450 standard; DNA; 17 AAA25450,

BP.

(first entry) 19-JUL-2000 AAA25450;

Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1948.

Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; haurpin ribozyme; antisense oligonucleotide; anticancer; breast cancer; endometrium cancer; ss.

Homo sapiens.

WO9954459-A2

28-OCT-1999.

99WO-US008547. 19-APR-1999; 98US-0082404P. 98US-00103636. 20-APR-1998; 23-JUN-1998;

(RIBO-) RIBOZYME PHARM INC.

Bellon L; Beigelman L, Mcswiggen JA, Karpeisky A, Zwick M, Jarvis T, Woolf T, Haeberli P; Matulic-Adamic J; Thompson JD, Reynolds M, 

WPI; 2000-013248/01.

nucleic acids that interact, and optionally cleave, target sequences, used to treat cancer.

Claim 77; Page 79; 148pp; English.

with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, The present invention describes nucleic acids (A) that interact stably

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This invention describes a novel retroviral expression vector (A) containing DNA sequences (I) for packaging vector RNA and for cellspecific expression of proteins or peptides encoding by heterologous DNA (II). The sequences controlling cell-specific expression contain a cellspecifically regulatable promoter region (P) from a human endogenous retrovirus (HERV) DNA sequence. The invention also describes (a) mRNA and NA of (A); (b) prokaryotic and eukaryotic cells containing (A) in integrated form; (d) virions containing a retroviral expression vector RNA derived from (A); (e) a method for producing the virions of (d); (f) a method for incorporating protein-encoding nucleic acid sequences into a eukaryotic cell by infection with the virions of (d); (f) a method for protein producing a packaging cell line, that contains at least one recombinant) retrovirus construct that contains at least one recombinant) retrovirus construct that contains at least one proteins of (A). (A) are used for cell- or tissue-specific expression of
                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Retroviral expression vector, useful in gene therapy, contains a promoter from a human endogenous retrovirus to provide cell-specific expression.
particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent cestrogen receptor hammerhead ribozyme sequences, and AAA25993 to AAA25905 represent their corresponding target sequences. AAA2593 to AAA26105 represent coestrogen receptor hairpin ribozyme sequences. AAA2619 to AAA26218 represent their corresponding target sequences. AAA26219 to AAA26271 represent other ribozyme sequences and antisense oligonucleotides used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cell-specific expression; tissue-specific expression; gene therapy; LTR; U3-R segment; long terminal repeat; retroviral expression vector;
                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                 DB 1; Length 17;
9.7e+02;
hes 0; Indels
                                                                                                                                                                                                                                                            Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                   0.6%; Score 17;
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 27; 67pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA98232 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                  Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; 88.
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Matches
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foreign genes for gene therapy and to produce virions for introducing (II) into the chromosomal DNA of eukaryotic cells, preferably mammalian and specifically human. (A) retain the advantages of usual retroviral promoters with all the signal structures required for transcription in a small region within the U3-R segment, but without their disadvantages (excessive strength and limited cell specificity). Since (A) are derived from endogenous (harmless) viral sequences, they do not introduce any new viral sequences into the genome and recombination will not create new types of retrovirus. The promoters provide cell or tissue specific expression, according to which HERV they are derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is that of a phosphorothicate oligonucleotide containing 20 T nucleobases, each having a 2'-methoxyethoxy group on its 5' ribosyl sugar moiety. It is an example of an oligomeric compound produced according to the methods of the invention. The invention provides compounds and methods for the preparation of mixed backbone oligomeric, or chimeric, compounds having phosphodiester internucleoside linkages in addition to phosphorothicate and/or phosphoramidate internucleoside linkages. The methods also include incorporation of boranophosphate internucleoside linkages. The methods utilise H-
                                                                                                                                                                                                                                                                                  Gaps
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/note= "phosphorothioate internucleoside linkages"
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Phosphorothioate oligonucleotide; H-phosphonate chemistry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag= a
/note= "2'-methoxyethoxy modified thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2'-Methoxyethoxy-modified phosphorothioate oligonucleotide.
                                                                                                                                                                                                                                                Length 17;
                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                         Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                              0.6%; Score 17; DB 1; Le
100.0%; Pred. No. 9.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 12; Page 34; 49pp; English.
                                                                                                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA50197 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 17; Conservative
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modified_base
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                                                                                                                                                                                                                                                    Query Match
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phosphonate intermediates that are coupled together forming contiguous regions of 1 or more H-phosphonate internucleoside linkages. Bach contiguous region is subsequently oxidized to phosphodiester, phosphorothioate, phosphoroamidate or boranophosphate internucleoside linkages prior to further elongation. Mixed backbone oligomeric compounds are prepared in this manner by oxidizing adjacent regions with different reagents. Oligomeric compounds of the invention are prepared using novel oxidation steps that oxidize a region of 1 or more H-phosphonate internucleoside linkages without degrading existing linkages that have been previously oxidized. The oligonucleotides obtained are useful as primers in PCR, probes, linkers, gene fragments and for other diagnostic tests on e.g. biological tissue, fluid, cells etc., as research reagents, and as antiviral agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive nucleotides from the 17 mer sequence, a sequence with, after optimal alignment, at least 80 % identity to the 17 mer sequence, a sequence that hybridizes to them under highly stringent conditions, or the complement of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, identifying, quantifying and/or amplifying a nucleic acid, e.g. as one component of a gene chib, in vitro as (antil sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, polypeptides, vectors containing the nucleic acids, cells containing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tumour suppression related human fukutin oligo SEQ ID No 352.
                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.6%; Score 17; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 9.7e+02; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzhaimer's disease and achizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for disgnosis and/or prognosis of these diseases. The polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
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/note= "Bases 9 and 10 are linked by a butanediol linker
which is represented as B in page 49 and X in page 59,
Fig 9 and 10 of the specification"
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                                                                                                                                                                                                                                                                                                      Gaps
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100.0%; Pred. No. 9.7e+02;
Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                 Local Similarity 100.
ses 17; Conservative
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to prevent or treat a disease characterised by the expression of a particular target RNA. The invention is used in gene therapy. The present sequence is an antisense oligo used to elicit human RNase (ribonuclease) H degradation of target RNA. This sequence is used in the exemplification of the invention
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//rotee Bases 9 and 10 are linked by a butanediol linker
which is represented as B in page 49 and Fig 5 and as X
in page 52, 55 and Fig 6 of the specification"
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                                                                                                                                         0.6%; Score 17; DB 100.0%; Pred. No. 9.7; ive 0; Mismatches
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Best Local Similarity 100.
Matches 17; Conservative
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They are useful for selectively preventing gene expression in a system. They are useful for selectively preventing gene expression in a sequence-specific manner, for hybridiaing to complementary RNA such as cellular RNA or viral RNA, to hybridiaing to complementary RNA. They are also useful therspeutically in formulations or medicaments to prevent or traget RNA. They are also useful therspeutically in formulations or medicaments to prevent target RNA. The invention is used in gene therapy. The present sequence is an antisense oligo used to elicit human RNase (ribonuclease) H degradation of target RNA. This sequence is used in the exemplification
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sequence is an antisense oligo used to elicit human RNase (ribonuclease)
H degradation of target RNA. This sequence is used in the exemplification
of the invention
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                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                    2'F-ANA antisense oligo #4, to elicit RNase H degradation of
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                                                                                                             Length 17;
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                                                                        Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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                                                                                                       0.6%; Score 17; DB 1; Le
Local Similarity 100.0%; Pred. No. 9.7e+02;
nes 17; Conservative 0; Mismatches 0;
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2'F-ANA antisense oligo #2, to elicit RNase H degradation of target RNA.
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/note= "Bases 4 and 5 are linked by a butanediol linker
which is represented as B in page 49 and Fig 5 and as X
in page 55 and Fig 6 of the specification"
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                                                                                                                                                                                           Gaps
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                                                                                                                         Length 17;
                                                                                                                      Ouery Match 0.6%; Score 17; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 9.7e+02; Matches 17; Conservative 0; Mismatches 0; Indels
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                                                           Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mangos MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AADS6447 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                               17 AAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ...17
*tag=
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*tag=
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of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense; ss.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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of the invention

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9. .10
/*tag= b
//tote= "Bases 9 and 10 are linked by a secouridine linker
which is represented as S in page 49 and X in page 57 and
Fig 1, 2, 7 and 8 of the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                  2'F-ANA antisense oligo #5, to elicit RNase H degradation of target RNA.
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                                                                                                                                                                                                                                                                                                                 Acyclic linker; gene expression; gene therapy; ribonuclease; RNase H;
                                    Gaps
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                                  0; Indels
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0.6%; Score 17; DB 1; Le
100.0%; Pred. No. 9.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER
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                                                                 2709 AAAAAAAAAAAAA 2725
                                                                                                                                                                                   BP
                                                                                                  17 AAAAAAAAAAAAAA 1
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                                                                                                                                                                                 AAD56450 standard; DNA; 17
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                                    Conservative
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*tag=
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Query Match
Best Local Similarity
Matches 17; Conservat
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                                                                                                                                                                                                                                                                                                                                     antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_feature
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ACC52437;
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                                                                                                                          RESULT 1310
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ADL48642/
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Analysis of the expression of the nucleotides can be used for diagnosis and or prognession of the nucleotides and polypeptides can be used for diagnosis and or prognession of the nucleotides and polypeptides can be used for diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                            primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             also be used to screen for their specific interactive molecules, potentially useful for treating diseases associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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  DB 1; Length 17; 9.7e+02;
                                         Indels
                                                                                                                                                                                                                                                                                                                                                                  Tumour suppression/reversion associated nucleotide #532.
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  0.6%; Score 17; DB 100.0%; Pred. No. 9.7; ive 0; Mismatches
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                                                                                  2709 AAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                         ADB40209 standard; DNA; 17 BP.
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                                                                                                                        AAAAAAAAAAAAAA 1
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Query Match
Best Local Similarity 100.1
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                      (revised)
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04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis.
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                                                                                                                                                                                    RESULT 1309
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New nucleic acid sequences associated with tumor suppression, regression,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents an isolated nucleic acid sequence associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  apoptosis or virus resistance are useful to diagnose and treat viral disease, development of tumor cells and cell degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; tumour suppressor; antitumour; cytostatic; tumour suppression; tumour regression; apoptosis; virus resistance; diagnosis; cellular degeneration.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
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100.0%; Pred. No. 9.7e+02;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                        Human tumour suppressor sequence #1204.
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1229 GATCTCCGAGATACAGG 1245
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                                                    1 GATCTCCGAGATACAGG
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Matches 17; Conservative
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
or a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (FTGDR),
[KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
on nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or gliona), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), orbon's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
cxpression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                         Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid amplification; RNA transcription; RNA polymerase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                        Haeberli P, Mcswiggen J, Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.6%; Score 17; DB 1; Length 17;
100.0%; Pred. No. 9.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 4 A; 4 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 100.0%; Pred. No. 9.7
                                                                                                                                                                                                                                                                                                                                                                                                            Claim 59; SEQ ID NO 2175; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of an oligo dT17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2535 GGCCTTGTCCTCAGCCA 2551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI34488 standard; DNA; 17 BP.
                                                                                                                                                                                                                           05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                                  03-APR-2002; 2002WO-US010512.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGCCTTGTCCTCAGCCA 1
                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                        Chowrira B,
                                                                                                                                                                                                                                                                                                                                  WPI; 2003-058513/05.
                                                                                                                                              WO200281628-A2
                                                                                             substrate; ds
                                                                                                                     Unidentified.
                                                                                                                                                                        17-0CT-2002
                                                                                                                                                                                                                                                                                                        Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADI34488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1312
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The invention relates to amplifying for RNA in a sample comprises eliminating single-stranded oligonucleotide from the transcription sample. The method involves synthesizing single-stranded cDNA by incubating the sample RNA with reverse transcriptase and an oligonucleotide primer that primes synthesis in a direction toward 5 end of the RNA; converting the single-stranded cDNA into double-stranded cDNA to form a transcription sample containing a cDNA template; eliminating single-stranded oligonucleotide from the transcription sample; and transcribing the cDNA template into RNA using an RNA polymerase. The method is useful for improving RNA polymerase based RNA transcription from a polymucleotide template. The method inhibits the undesired non-femplate derived production of RNA in the transcription reaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Intramolecular base pair; intermolecular base pair; unstructured nucleic acid; UNA; molecular biology; nucleic acid chemistry; polymerse extension reaction; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / Match 0.6%; Score 17; DB 1; Length 17; Local Similarity 100.0%; Pred. No. 9.7e+02; nes 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                          31-MAY-2002; 2002US-0384454P.
                                                                                                                                                    30-MAY-2003; 2003WO-US017103
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31-JUL-2000; 2000US-00632639.
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                                                                                                                                                                                                                                                                                                (JANC ) JANSSEN PHARM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transcription reaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-035466/03.
                                                                                                                                                                                                                                                                                                                                                                               Zhu JY;
WO2003102243-A1.
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                                                                          11-DEC-2003.
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WPI; 2004-487902/46
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                      ADP86137;
                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                          The present invention provides a system for the production of nucleic acids with reduced levels of intramolecular base pairing (secondary structured) and intermolecular base pairing by generating unstructured nucleic acids (UNAs). The invention is useful for generating nucleic acid having a reduced ability to hybridise. The invention is also useful in molecular biology and nucleic acid chemistry. The present sequence is an annealing primer used to generate single-stranded labelled unstructured nucleic acid (UNA) by polymerase extension reaction (PCR). This sequence is used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunostimulatory oligonucleotide; immune response; allergy; asthma;
                                                                                Generating nucleic acid having reduced ability to hybridize for use in molecular biology, comprises providing nucleotide triphosphates to synthesize nucleic acid complementary to a template nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         viral infection; bacterial infection; cancer; lymphoma;
intraepithelial neoplasm; melanoma; neuroblastoma; Hodgkin's lymphoma;
carcinoma; sarcoma; gene therapy; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                      Sequence 17 BP; 17 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           CpG immunostimulatory oligonucleotide #49
                                                                                                                           Disclosure; SEQ ID NO 16; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    base= OTHER
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                                         Wolber P;
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25-SEP-2003; 2003US-0506108P.
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                                                                                                                                                                                                                                                                                                                                                                                             ADP86178 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                        Sampson JR, Ach RA,
                                                            WPI; 2004-364526/34.
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SAMPSON J R.
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         ACH R A.
WOLBER P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                  ADP86178;
        (ACHR/)
(WOLB/)
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New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CpG immunostimulatory oligonucleotide #8.
                                                                                                                                                               Example; SEQ ID NO 49; 104pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunostimulatory oligonucleotide.
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25-SEP-2003; 2003US-0506108P.
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                                                                                        cancer, cervical cancer.
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The invention relates to a class of CpG immunostimulatory
oligonucleotides containing a 5'TCG motif or a CG at or the 5' end that
are useful for stimulating an immune response. Oligonucleotides and
compositions of the invention are useful for treating allergy or asthma,
viral and bacterial infections and cancer e.g. biliary tract cancer,
breast cancer, cervical cancer, choriocarcinoma, colon cancer,
condomerrial cancer, gastric cancer, lymphomas, intraepithelial neoplasms,
liver cancer, lung cancer (e.g. small cell and non-small cell), melanoma,
cutroblastomas, ovarian cancer, pancreatic cancer, prostate cancer,
rectal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer,
cetal cancer, connective tissue cancer, osesophageal cancer, brain
and CNS cancer, connective tissue cancer, osesophageal cancer, ere
chodykin's lymphoma, larynx cancer, oral cavity cancer, skin cancer,
testicular cancer, as well as other carcinomas and sarcomas. The
invention is also useful in gene therapy. The present sequence is a CpG
immunostimulatory oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fires a hosphorylated poly dT, which binds to a bridging oligonucleotide"
New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast
                                                                                                                                                                                                                                                                                                                                                                                           0.6%; Score 17; DB 1; Length 17; 100.0%; Pred. No. 9.7e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA amplification; hybridization; primer; ss.
                                                               Example; SEQ ID NO 8; 104pp; English.
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2002US-0367438P.
2003US-00393519.
2003WO-US009232.
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2000US-0234060P.
2001US-0261231P.
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2001US-00908950.
2001WO-US029589.
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2004US-00825776.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Poly dT primer SEQ ID NO:3
                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match
Best Local Similarity 100.
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-NOV-2005 (first entry)
                                    cancer, cervical cancer.
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25-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEC07079;
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Gaps ö

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The invention relates to a method for determining the presence of at least one specific nucleotide sequence in a target nucleic acid extracted from a biological sample comprises preparing a primer oligonucleotide comprising a first bridging sequence with a primer portion composed of a terminating and group attached to one end of the first bridging sequence and a terminating end group attached to the other end. The methods and compositions of the present invention are useful for assaying the presence of specific nucleic acids sequences, particularly for hybridizing nucleic acids in solutions or on surfaces like microarrays, and their amplification using RNA polymerase. The present sequence represents a poly dT primer, which is used in the exemplification of the
                                                                                                                                                                                                                                                                                       Assaying the presence of specific nucleic acids sequences using primer oligonucleotides having a first bridging sequence with a primer portion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
0.6%; Score 17; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 9.7e+02;
Matches 17; Conservative 0; Mismatches 0;
                                                                                                                                                                      Howerton
                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 3; 13pp; English.
                                                                                                                                                                   Schwalm J,
                                                                                                                                                                                                                                                                                                                                                              composed of random nucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-2005; 2005WO-US011827.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .285/c
AED81285 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                               Kadushin J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vollmer J;
GETTS R C.
KADUSHIN J.
SCHWALM J.
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                                                                                                                                                                                                                             WPI; 2005-618097/63
                                                                                                  HOWERTON K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present invention.
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                                                                                                                                                                   Getts RC,
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                                                                                                  (HOWE/)
                                                                (SCHM/)
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C directeotide, and nt and nt and nt are polymucleotides that do not include directed, where NI does not include 5' Z nucleotide, and where a 3' polymucleotide consisting of the YZ dinucleotide and the NO shall be polymucleotide consisting of the YZ dinucleotide and where a 3' polymucleotide consisting a number of nucleotides in the oligonucleotide; and at most 45¢ of the number of nucleotides in the oligonucleotide; and and if a denotes the 3' can do fet he oligonucleotide, and where X is a T or modified T nucleotide, YI is a T or modified G nucleotide, where NI does not include a C or modified of 5-10 nucleotides, where NI does not include a C dinucleotide of 5-10 nucleotides, where NI does not include a C dinucleotide of 5-10 nucleotides. Also described are: (1) a polymucleotide of 5-10 nucleotides. Also described are: (1) a c polymucleotide of 5-10 nucleotides. Also described are: (1) a c polymucleotide of 5-10 nucleotides. Also described are: (1) a c method of pectifically increasing interleutin (IL)-10 expression in the atherapeutic agents, monoclonal antibodies, and anticancer agents; radiotherapeutic agents, monoclonal antibodies, and anticancer agents; c relative to interferon (IRN)-alpha expression in a subject. compression (IRN)-apha expression in a subject composition to the subject of inducing an antigen. Also method of inducing an antigen and administering an immunostimulatory nucleic acid or composition is administering an immunost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigenspecific response to the implant in the subject. The oligonucleotide includes at least I modified internucleotide linkage such as a phosphorothioate linkage. The oligonucleotide, methods and compositions of the invention are useful for treating allergies, asthma, autoimmune diseases, arthritis, systemic lupus erythemarosus, multiple solerosis, crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune neuropathies such as Guillain-Barre, autoimmune uvaitis, autoimmune hemolytic anemia, pernicious anemia, autoimmune thrombocytopenia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Crohn's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma,
                                                                                                                 XYNIYZN2 3' where 5' denotes the 5' end of the oligonucleotide and 3' denotes the 3' end of the oligonucleotide, where X is a T or modified T nucleotide, Y is a C or modified C nucleotide, Z is a G or modified G nucleotide, and NI and N2 are polymucleotides that do not include a CG nucleotide, and NI and N2 are polymucleotides that do not include a CG
                                                                                     invention relates to an oligonucleotide having the formula: (a) 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, villigo, Crob Disease, ulcerative colitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated diabetes mellitus, Grave's Disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oligonuclectide used in experiments in the examples of the present
                            Example; SEQ ID NO 43; 111pp; English
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temporal arteritis, anti-phospholipi
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sambles.

0.6%; Score 17; DB 1; Length 17; 00.0%; Pred. No. 9.7e+02; Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other; 100.0%; Pred. No. 9.7 Query Match Best Local Similarity 100.0 Matches 17; Conservative

17 AAAAAAAAAAAAAA 1

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RESULT 1318

**AEF82502** 

Novel 18S ribosome RNA gene derived from common marmoset, or its partial fragment, useful as internal standard for measuring difference in expression level of gene of interest in two or more types of test 18s ribosomal RNA; 18s rRNA; RNA detection; DNA detection; expression; Common marmoset 18S ribosome PCR primer SEQ ID NO:4. AEF82502 standard; DNA; 17 BP 09-JUL-2004; 2004JP-00202891. 04-MAR-2005; 2005JP-00060329. CO LTD 20-APR-2006 (first entry) (SUMO ) SUMITOMO CHEM WPI; 2006-150097/16. Yamada T, Oeda K; ss; PCR; primer. JP2006042804-A. 16-FEB-2006. Synthetic. AEF82502; 

The invention relates to a novel 18S ribosome RNA gene (I) derived from a common marmoset, or its partial fragment (AEF82499). Also claimed is a composition for detecting DNA or RNA, comprising the 18S rRNA gene. The 18S rRNA gene is useful as an internal standard or a reference of the expression level of agene in the test sample during the measurement of the difference in the expression level of a gene of interest in two or more types of test sample, based on the difference in the transcription product amount of the gene, where the test sample is derived from a common marmoset. The transcription product amount of the gene of interest PCR. The present sequence represents a PCR primer used in the invention to synthesise 18S rRNA cDNA. transcriptaseis measured using a DNA array or by quantitative reverse Disclosure; SEQ ID NO 4; 21pp; Japanese.

Sequence 17 BP; 0 A; 0 C; 0 G; 17 T; 0 U; 0 Other;

Gaps ö Score 17; DB 1; Length 17; Pred. No. 9.7e+02; 0; Indels Query Match Best Local Similarity 100.0%; Fred. No. 9.76 Gongervative 0; Mismatches

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2709 AAAAAAAAAAAAAA 2725 17 AAAAAAAAAAAAAAA

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AAT94668 standard; DNA; 18 BP. 27-MAR-1998 (first entry) AAT94668; AAT94668 \*\*\*\*

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Gaps

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Indels

RESULT 1319

Anchored poly(T) oligonucleotide polyT-AnchC.

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2709 AAAAAAAAAAAAAA 2725

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Sakaki Y;
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                                                                                                                                                                                                                                                   RESULT 1321
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                                                                                                                                                                                                                    Anchored poly(T) oligonucleotides polyT-anchA (AAT94667), polyT-anchC (AAT94668) and polyT-anchG (AAT94669) are complementary to the upstream region of a polyadenylation sequence. They were used to prime CDNA synthesis from snapdragon (Antirrhinum majus) petal and leaf RNA, and were also utilised in the PCR amplification of plant cytochrome P450 sequences (see also AAT94670-73). A cDNA clone (see AAT94657) encoding flavonoid 3' hydroxylase (see AAN95704) was isolated using a differential display approach. This can be used to manipulate the pigmentation of
Flavonoid 3' hydroxylase; pigmentation; flower colour; transgenic plant; snapdragon; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Flavonoid 3' hydroxylase; pigmentation; flower colour; transgenic plant;
                                                                                                                                                                          Novel flavonoid 3'-hydroxylase(s) from flowering plants - and corresponding DNA, used in the manipulation of pigmentation in plants.
                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                   Length 18;
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                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 0 A; 1 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 0.6%; Score 17; DB 1;
100.0%; Pred. No. 1e+03;
iive 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anchored poly(T) oligonucleotide polyT-AnchG.
                                                                                                                                       Michael MZ;
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                                                                                                                                                                                                       Example 15; Page 59; 234pp; English.
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                                                                                 97WO-AU000124.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT94669 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                      Holton TA,
                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     snapdragon; primer; ss.
                                                                                                                    (FLOR-) FLORIGENE LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (FLOR-) FLORIGENE LTD
                                                                                                                                                        WPI; 1997-448691/41.
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                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                              transgenic plants
                                                                                                                                       Brugliera F,
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                                            WO9732023-A1
                                                                                28-FEB-1997;
                                                                                                  01-MAR-1996;
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                                                              04-SEP-1997
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                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT94669;
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                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                         (AAT94668) and polyT-anchG (AAT94669) are complementary to the upstream region of a polytaderylation sequence. They were used to prime cDNA synthesis from snapdragon (Antirrhinum majus) petal and leaf RNA, and were also utilised in the PCR amplification of plant cytochrome P450 were slso utilised in the PCR amplification of plant cytochrome P450 aguences (see also AAT94670-73). A cDNA clone (see AAT94657) encoding flavonoid 3' hydroxylase (see AAN95704) was isolated using a differential display approach. This can be used to manipulate the pigmentation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel apoptosis-related DNAs and proteins - for diagnosis, preventing or treating diseases associated with apoptosis.
Novel flavonoid 3'-hydroxylase(s) from flowering plants - and corresponding DNA, used in the manipulation of pigmentation in plants.
                                                                                                                                                                         poly(T) oligonucleotides polyT-anchA (AAT94667), polyT-anchC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17; DB 1; Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. No. 1e+03;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 0 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treatment of apoptosis-related diseases
                                                                                                        Example 15; Page 59; 234pp; English,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 49; 70pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleotide sequence PCR primer 7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV54170 standard; cDNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-495844/42,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           transgenic plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9839437-A1
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AAV37712;

RESULT 1322

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AAV37712

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Determining the nucleotide sequence of a nucleic acid analyte - using
                                                                                                                                                                                                                                                           /*tag= a
/note= "phosphorothioate internucleotide linkages"
                                                                                                                              phosphorothioate; electrospray ionisation-Fourier transform; mass spectrometry; off-resonance excitation; ss.
                                                                                                  Phosphorothioate oligodeoxynucleotide.
                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Fig 3A; 25pp; English.
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AAV07750 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAY-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    electro-spray ionisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 100.
hes 17; Conservative
                                                                                                                                                                                                                                        ...18
                                                                                                                                                                                                                                                                                                                                                                                                                                              (HYBR-) HYBRIDON INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-520830/44.
                                                                                                                                                                                                                                        misc_difference
                                                                                                                                                                                                                                                                                                                                                                             12-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                               14-MAR-1997;
                                                                                                                                                                                                                                                                                                           WO9840520-A1
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                                                                  02-DEC-1998
                                                                                                                                                                                       Synthetic
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                                 AAV07750;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang BH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1324
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Matches
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ID AAX1
 XXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes novel proteins isolated from cDNA clones. AE648 11; AE693 11; AK438 11; AK609 11; AM1060 11; AQ2 11; K433 11; or 1256 11, deposited as ATCC 98237. The present sequence represents the 3'-portion of AQ2 11 isolated from a human ovary cDNA library. The proteins from the present invention may be administered in a composition to prevent, treat or ameliorate a medical condition. The proteins may exhibit biological activities such as nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating or suppressing activity, haematopoiesis regulating activity, tissue growth haemostatic and thrombotic activity, chemicaticiferentiation activity, haematory activity, cadherin/tumour invasion suppressor activity, until inflammatory activity, cadherin/tumour invasion suppressor activity, tumour inhibition activity and other activities. (Updated on 25-MAR-2003
                                                                                                                                                                                                                                                                                        Human; secreted protein; murine adult spleen; human foetal kidney; ovary;
bone marrow; thymus; AE648_11; AE693_11; AK438_11; AK609_11; AM1060_11;
AQ2_11; K433_11; L256_11; prevent; treat; ameliorate; medical; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New secreted proteins and associated polynucleotides - obtained from murine adult spleen, human foetal kidney, human ovary, murine bone marrow and murine adult thymus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6%; Score 17; DB 1; Length 18; 0.0%; Pred. No. 1e+03; 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 17 A; 0 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                           Human protein AQ2_1i 3'-portion and polyA tail.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lavallie ER, Racie LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.6*; Score 1,; C. Best Local Similarity 100.0%; Pred. No. 1e+ Matches 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 58; 75pp; English.
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                   2707 CTAAAAAAAAAAAAA 2723
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97US-00960024
                                                                                                                                         AAV37712 standard; cDNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jacobs K, Mccoy JM, Lava
Spaulding V, Agostino MJ;
                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to correct PR field.)
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                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                   WO9820130-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1996;
29-OCT-1997;
                                                                                                                                                                                                         25-MAR-2003
07-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAY-1998
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The invention relates to an analytical method for determining the nucleotide sequence of nucleic acid analytes, including chemically nucleotide sequence of nucleic acid analytes, including chemically modified oligonucleotides. This new method utilises electrospray constants. The ions are excited by sustained off-resonance excitation with single shot excitation, and the target fragmented by collisionally activated dissociation by a neutral cast, e.g. carbon dioxide. Alternatively, the excitation and dissociation can be nozale skimmer dissociation. The method is used in molecular biology and biomedical applications. The method is used in molecular can be nozale skimmer dissociations. The method, utilising electrospray is extremely rapid and acts directly on the oligonucleotide. The method is effective for a variety of nucleic acid analytes, particularly successfully sequenced. The present sequence represents a successfully sequenced. The present sequence represents a proposition of the present sequence represents and successfully sequenced.
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Pred. No. 1e+03;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 17 A; 0 C; 0 G; 1 T; 0 U; 0 Other;
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8 셤 RESULT 1323

AAV07750

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RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.
                                                                                                                                                                                                                                                       Score 17; DB 1; Length 18;
Pred. No. 1e+03;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                        RT-PCR primer of the invention SEQ ID 13,
                                                                                                                                     Disclosure; Page 11; 19pp; Japanese
                                                                                                                                                                                                                                                      0.0.,
100.0%; Pre-
                                                                                                                                                                                                                                                                                  2708 TAAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                               AAX18372 standard; DNA; 18 BP.
                                                            97JP-00208312.
                                                                         97JP-00208312
                                                                                                                                                                                                                                                                                                17 TAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                             97JP-00208312.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          97JP-00208312
                                                                                      (TAKI ) TAKARA SHUZO CO LTD
                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                     17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TAKI ) TAKARA SHUZO CO
                                                                                                    WPI; 1999-183822/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-183822/16.
                                                                                                                                                                                                                                                              Best Local Similarity
                                JP11032765-A
                                                           18-JUL-1997;
                                                                         18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                            11-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                 JP11032765-A
                                                                                                                                                                                                                                                                                                                                                                                                                                            18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JUL-1997;
                                              09-FEB-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                               09-FEB-1999
                    Synthetic.
                                                                                                                                                                                                                            sednences
                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                        Query Match
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This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta a N3'; or (X)m5'-(gamma)** (A) the sequence of a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine, or cytosine; N = adenine, guanine or thymine; and a thymine; amount a thymine; amount a thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptides having at least two new nucleotides - useful as primers in RT-
                       88.
RT-PCR primer; DNA sequence determination; gene sequence analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18 BP; 1 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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Gaps

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Score 17; DB 1; Length 18; Pred. No. 1e+03; 0; Mismatches 0; Indels

0.6%; Scor. 100.0%; Pred. No. 10.

15; Conservative

Query Match Best Local S

Best Loca Matches

2708 TAAAAAAAAAAAA 2724

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17

BP.

AAA40563 standard; cDNA; 18

RESULT 1326

AAA40563

(first entry)

16-NOV-2000

Gaps

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AAA40563;

Sequence 18 BP; 2 A; 0 C; 0 G; 16 T; 0 U; 0 Other;

sednences

This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta a N3', or (X)m5'-(gamma) K-delta-N3'; where X = a labelled compound and/or nucleotide with voluntary sequence; m = 0 or 1; alpha = thywine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, quanine or cytosine; N = adenine, guanine, cytosine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, quanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene

Disclosure, Page 11; 19pp, Japanese.

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antiviral; immunosuppressive; antiinflammatory; vulnerary; cytokine; cell proliferation; differentiation; regulator; treatment; tumor; autoimmune disease; inflammatory disorder; wound; microbial infection; viral disease; graft versus host reaction suppression; ss.
                                                                                                                                                                                                                        ပဲ
                                                            cytostatic; immunostimulatory; antimicrobial;
                                                                                                                                                                                                                        Evans
                                                                                                                                                                                                                       Collins-Racie LA,
                                          Human adult ovary cDNA fragment AQ2_1i #2.
                                                                                                                                                                                                                       Lavallie ER,
                                                                                                                                                                                                                                Bowman MR;
                                                                                                                                                                   99WO-US031005.
                                                                                                                                                                                    98US-00220876.
                                                                                                                                                                                                     (GEMY ) GENETICS INST INC.
                                                                                                                                                                                                                      Mccoy JM, 1
Treacy M,
                                                           Secreted protein;
                                                                                                                                WO200037630-A1.
                                                                                                                 Homo sapiens
                                                                                                                                                                   22-DEC-1999;
                                                                                                                                                                                    23-DEC-1998;
                                                                                                                                                                                                                      Jacobs K, M
Merberg D,
                                                                                                                                                  29-JUN-2000
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Secreted human proteins AS296-11 and AS34-11, useful for treating tumors, autoimmune diseases, inflammatory disorders, wounds, microbial infections

WPI; 2000-442661/38.

P-PSDB; AAB10274

Disclosure; Page 269; 293pp; English.

and viral diseases.

This invention describes novel secreted human proteins (I) which have cytostatic, immunostimulatory, antimicrobial, antiviral, immunosuppressive, antimilammatory and vulnerary activity and which act as cytokine, cell proliferation or differentiation regulators. (I) is useful for treating tumors, autoimmune diseases, inflammatory disorders,

Peptides having at least two new nucleotides - useful as primers in RT-

ngs.res

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21-APR-2005
                                                     Unidentified
                                                                                                                                        04-OCT-1999;
                                                                          US6271002-B1
                                                                                                                  04-OCT-1999;
                                                                                               07-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADX69542;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to identification of genes and proteins of adipose tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperunicemia and sleep apnea syndrome. The genes (AAZ90631-633) and the proteins (AAX97598-Y67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose tissue genes
wounds, microbial infections and viral diseases. (I) is also useful for suppressing graft versus host reaction. AAA40490-A40580 represent cDNA fragments that encode the secreted proteins AAB10226-B10288 described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A physiologically active protein specifically derived from mammal tissue
                                                                                                                                                                                                                                                                             Adipose tissue; obesity; diabetes; hyperlipemia; hypertension; human; arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss
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                                                                                                Gaps
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                                                                                                ;
0
                                                                        Query Match

0.6%; Score 17; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 1e+03;
Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.6%; Score 17; DB 1; Length 18;
100.0%; Pred. No. 1e+03;
1ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 BP; 1 A; 0 C; 2 G; 15 T; 0 U; 0 Other;
                                                     Sequence 18 BP; 17 A; 0 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                           Human adipose tissue gene amplifying primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 18; 50pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.08; FIL
                                                                                                                    2709 AAAAAAAAAAAAAA 2725
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                                                                                                                                         AAAAAAAAAAAAA 18
                                                                                                                                                                                   AAZ90640/c
ID AAZ90640 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                    (NISB ) JAPAN TOBACCO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                    JP2000037190-A.
                                                                                                                                                                                                                                                                                                                                                                           23-JUL-1998;
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                                                                                                                                                                                                                                      13-JUN-2000
                                                                                                                                                                                                                                                                                                                Homo sapiens
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The invention relates to methods and kits for amplification of mRNA using a primer in PCR that contains an RNA polymerase (RNAP) promoter. The invention provides methods for amplification and detection of RNA derived from a population of cells, preferably eukaryotic cells and most preferably mammalian cells, which methods preserve fidelity with respect to sequence and transcript representation and additionally enable amplification of extremely small amounts of mRNA. The method and kit are useful for amplifying and detecting RNA derived from a population of cells, especially eukaryotic cells like mammals. The RNAS generated are useful for profiling gene expression in different populations of cells. The present sequence is a mRNA fragment used in 3' end PCR/IVT (in vitro
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                                                                  RNA polymerase; RNAP; RNA detection; IVT; in vitro transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amplifying and detecting RNA derived from a population of cells Lemploying a primer that contains an RNA polymerase promoter in a polymerase chain reaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO 11
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mRNA fragment used in 3' end PCR/IVT method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.6%; Score 17; DB 1; Length 18; 100.0%; Pred. No. 1e+03; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 17 A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; PCR; primer; transformation; plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transcription) method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ROSE-) ROSETTA INPHARMATICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Fig 1; 29pp; English.
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                                                                                                                                                                                                                                                                                                                                                              99US-00411074.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-OCT-2002; 2002KR-00064822
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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hes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Linsley PS, Schelter JM;
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Preparing transformed monocotyledon rice, with t-DNA tagging vector comprising enhancer element for activation tagging and reporter gene for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of ababled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                 monocotyledons. The present sequence represents a Monocotyledon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                           , DB 1; Le...
'9, 1e+03;
0; Indels
                                                                                                                                                                                The invention relates to a method of preparing transformed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.6%; Score 17; DB 1; Length 19; 100.0%; Pred. No. 1e+03; Astive 0; Mismatches 0; Indels
                               Mun SO
                                                                                                                                                                                                                                           Sequence 18 BP; 1 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
0; Mismatches
                               HG,
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                               Kang
                                                                                                                                                Disclosure; SEQ ID NO 11; 17pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            digestion with restriction enzymes.
                                                                                                                                                                                                             transformation associated PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 5; 11pp; Japanese.
                             Jung DH,
                                                                                                                                                                                                                                                                                                                                     2707 CTAAAAAAAAAAAA 2723
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ75558 standard; DNA; 19 BP
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es 17; Conservative
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Best Local Similarity 100.
Matches 17; Conservative
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                                                          WPI; 2004-589679/57
                               An GH,
                                                                                                                  gene trapping.
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Matches
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Human, antisense, bronchoconstriction, allergy, hyposecretion, pain, respiratory tract inflammation, adenosine sensitivity; lung; cancer; surfactant depletion, antiallergic, antiinflammatory; antiasthematic, analgesic, hypotensive, immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                              expression; reverse transcription; primer; cDNA;
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                                                                                                                                                                                                           Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 17; DB 1; Length 19;
100.0%; Pred. No. 1e+03;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 0 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AI095492-derived oligonucleotide SEQ ID 3936.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure, Page 5, 11pp; Japanese.
                                                                                                                                                                                                                                                            aggregate; restriction enzyme; ss.
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2709 AAAAAAAAAAAAAA
                                                                                                              AAQ75550 standard; DNA; 19
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                                                                                                                                                                                                                                           Analysis; gene
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                                                                                                                                                                                                                                                                                             Synthetic.
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                                                                               RESULT 1331
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Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.

Reverse transcription primer used in cDNA analysis technique.

(first entry)

04-AUG-1995

AAQ75574;

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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating comprising oligonuclectides, effective for alleviating comprising oligonuclectides, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, configuration or thyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regularing expression of a target polypeptide associated with lung airway or lung expression or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thy wildines present in the target RNA serves to prevent the breakdom of the oligonuclectides into products that free adenosine into the system content of the adenosine content of the adenosine content of the adenosine content of the adenosine content of the endiance adenosine content of the endiance adenosine of the content of the target RNA se
respiratory distress syndrome, allergic rhinitis, pulmonary hypertension, emphysema; chronic obstructive pulmonary disease, cancer, bronchitis; pulmonary transplantation rejection, ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotide containing less percentage of adenosine, targeted uncleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     composition for treating asthma, has antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Leus
o. 1e+03;
0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 16 A; 0 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 3936; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .6%; Score 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sandrasagra A, Ka
,, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286036P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bronchodilating agent.
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Tang L,
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                       WO200285309-A2
                                                                                                                                                                         Homo sapiens
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Miller S,
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Gaps

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0.6%; Score 17; DB 1; Length 20; 100.0%; Pred. No. 1e+03; tive 0; Mismatches 0; Indels

2709 AAAAAAAAAAAAAA 2725

17; Conservative

Matches

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Local Similarity

Query Match

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BP.

AAQ75605 standard; DNA; 20

RESULT 1334

AAQ75605

Seguence 20 BP; 0 A; 2 C; 1 G; 17 T; 0 U; 0 Other;

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of ababled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily

expression - by amplification of mRNA followed

by digestion with restriction enzymes. Disclosure; Page 5; 11pp; Japanese.

of cDNA and gene

Analysis

WPI; 1995-018287/03.

(NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.

93JP-00112515. 93JP-00112515.

16-APR-1993; 16-APR-1993;

01-NOV-1994.

JP06303997-A.

Synthetic.

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Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                           Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                    (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                93JP-00112515.
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                          (first entry)
                          04-AUG-1995
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                                                                                           Synthetic.
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Mismatches

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Conservative

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ઠ 유 AAQ75574 standard; DNA; 20 BP

RESULT 1333 AAQ75574,

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method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                 Sequence 20 BP; 0 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
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                                  by digestion with restriction enzymes.
                                                  Disclosure; Page 5; 11pp; Japanese
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aggregate; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                 AAQ75572 standard; DNA; 20 BP.
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEC files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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AAQ75604 standard; DNA; 20
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Query Match Best Local Similarity

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and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                               gene expression - by amplification of mRNA followed
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                                    Reverse transcription primer used in cDNA analysis technique.
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Or Indels
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genetic engineering; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 1 C; 2 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                          (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                       digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of a PCR primer #1
                                                                                                                                                                                                                                       Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                     0.6%; Scor
100.0%; Pred
                                                               restriction enzyme; ss
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Best Local Similarity 100.0%;
Matches 17; Conservative.
                   (first entry)
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(TAKA/) TAKAHASHI K.
                                                                                                                                                                                                               Analysis of cDNA and
                                                                                                                                                                                             WPI; 1995-018287/03.
                                                      Analysis; gene
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                                                               aggregate;
                                                                                 Synthetic.
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AAQ75571;
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                                                                (PCR) by using
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                                                         Carrying out a thermal cycle of polymerase chain reaction (PCR) by using a substrate on which a DNA is immobilized used in medical, biochemical, molecular biological and gene engineering fields.
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Pred. No. 1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 3 A; 0 C; 0 G; 17 T; 0 U; 0 Other;
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Pred. No. 1e+03;
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100.0%; Pred
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                                                                                                                                                                                             Example; Page 9; 13pp; Japanese
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WPI; 2002-630904/68.
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Best Local Similarity
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intiation retained to a plantameterial composition, which it is active agent comprising an oligimaterial composition, which is active agent comprising a polymentation codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 6' or 3' end genomic flanking regions, 6' or 3' end genomic flanking regions, 6' or antialized an encoding a polymeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an ential flammatory second active agent composition of the invention thas antialized and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or arespiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                  Human, antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiguinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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Mismatches
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Tang L, Shahabuddin S;
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Matches 17; Conservative
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Miller S,
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Length 20;

Score 17; DB 1; Pred. No. 1e+03;

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Query Match Best Local Similarity

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intran-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention thas antiinflammatory, antiallargic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gene therapy. The composition may have a cuse in antisense gratery and alignant corposition may have a crowenting a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entiinflammatory steroid in a subject, for reducing or depleting levels of ceceptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject is tissue, or treating bronchoconstriction, clung surfactant in a subject is tissue, or treating bronchoconstriction, or ung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed presented in the printed and the control of the co
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Pred. No.
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Tang L, Shahabuddin S;
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first active agent comprising an oligomorelectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nuclectides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction as decond active agent comprising an entilnflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallargic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entilnflammatory steroid in a subject, for reducing levels of contition, also of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or ung surfactent in a subject's tissue, or treating bronchoconstriction, lung surfactent in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allerajes, or a respiratory disease or condition. Note: The sequence date for this patent is not represented in the printed of the first into int, which was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                         Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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  Mismatches
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i, Shahabuddin S;
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Tang L,
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Length 20;

Score 17; DB 1; Pred. No. 1e+03;

0.6%; Score 17;

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Query Match Best Local Similarity

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation coodon, coding regaion, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antiasnes gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing bronchoconstriction, lung surfactant in a subject's tissue, or treating bronchoconstriction,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                             Human; antisense; lung dysfunction, nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodiation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lung allergies, or a respiratory disease or condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17; DB 1; Length 20;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 16 A; 2 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 4256; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Katz E,
S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sandrasagra A,
, Shahabuddin
2709 AAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                        Human oligonucleotide sequence.
                                                                                                                                      BP.
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                                                                                                                                    ABZ89014 standard; DNA; 20
                                     19 AAAAAAAAAAAAA
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                               RESULT 1346
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating conditions. Trespiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cauteding adenosine sensitivity, levels of adenosine (A) or (A) receptors, collgonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction also describes a kit, that comprises: (a) a delivery device, in separate containers (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antialflammatory, antiatementic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung or with a disease or condition also are associated condition and and proportion, and and an and an appropriate condition as applicant pression and an application.
                                                                                                                                                                                                                                                                                                                                                                                  respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                     antisense; bronchoconstriction; allergy; hyposecretion; pain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation, allergies, asthma, impeded respiration, respiratory
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Indels
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Mismatches
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Shahabuddin S;
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                                        2707 CTAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                  (first entry)
Matches 17; Conservative
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15;
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             transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antialflammatory; antiasthmatic; analgesic; hypotensive; immunosupptessive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
pulmonary
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chronic obstructive pulmonary disease,
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                                                                                                                                                                                        Length 20;
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                                                                                                                                                   Seguence 20 BP; 2 A; 1 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                        Score 17; DB 1;
Pred. No. 1e+03;
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Tang L, Shahabuddin S;
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100.0%; Pred
0; M
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                                                                                                                                                                                                                                                                                                                                                                                                     ABD25244 standard; DNA; 20
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                                                                                                                                                                                                                              17; Conservative
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                                                                                                                                                                                                                                                                   2709
                                                                                                                                                                                                                                                                                                                                                                                                                                          ABD25244;
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of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrencegic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered corposition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production of target polypeptide present in the lungs. The reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung of inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary disease, pulmonary cinflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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100.0%; Pred. No. 1e+03;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 16 A; 2 C; 0 G; 2 T; 0 U; 0 Other;
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Matches 17; Conservative
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20-MAY-2003; 2003WO-US016084. 20-MAY-2002; 2002US-0381857P. Crosby SD, Nalseth AE; (PHAA ) PHARMACIA CORP WPI; 2004-035034/03. WO2003099215-A2. Homo sapiens. 04-DEC-2003. ဌ Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and ä Pabalan J, Aguilar Katz E, Claim 15; SEQ ID NO 5138; 763pp; English. i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 23-APR-2002; 2002WO-US013143. (EPIG-) EPIGENESIS PHARM INC. 24-APR-2001; 2001US-0286036P. bronchodilating agent

WPI; 2003-093058/08.

Li Y,

Miller S,

vyce JW,

31-OCT-2002.

This invention describes a novel composition (a) a first active agent, comprising oligonuclecides, effective for alleviating bronchoonstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung or inflammation, allergies and/or surfactant hypoproduction are associated with a allergies and/or surfactant hypoproduction are associated with a allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, pulmonary transmation, allergies, asthma, impeded respiration; pulmonary transmation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary impeded respiration; pulmonary transmation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary incertions and represent in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine content of the anti-sommer and the prevent. In the print, heart, kidney envised to prevent the prevent. e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it

Sequence 20 BP; 18 A; 0 C; 2 G; 0 T; 0 U; 0 Other;

Gaps ö le+03; 0.6%; Score 17; DB 1; Length 20; 100.0%; Pred. .v.. 17; Conservative Local Similarity Query Match Matches

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ADH67409 standard; DNA; 20 BP (first entry) 25-MAR-2004 ADH67409;

RESULT 1350

ADH67409/

Human glucocorticoid receptor-specific antisense oligonucleotide #4243.

antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; fumour formation; diabetes; obssity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2 -methoxyethyl; 2 -MOE.

New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.

Claim 4; SEQ ID NO 4243; 985pp; English

to nucleic acids encoding a mammalian glucocorticoid receptor. The antiennes oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence repersents an antisense oligonucleotide that targets the human glucocorticoid receptor gene NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone. The invention comprises an antisense oligonucleotides that are targeted

Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;

Score 17; DB 1; Length 20; Pred. No. 1e+03; Query Match 0.6%; Score 17; DB Best Local Similarity 100.0%; Pred. No. 10+ Matches 17; Conservative 0; Mismatches 2709 AAAAAAAAAAAAAAA 2725 4 20 AAAAAAAAAAAAAAAA 셤 ઠ

ADK75123 standard; DNA; 20 BP (first entry) 20-MAY-2004 ADK75123 RESULT 1351 

Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; 88. Chimeric phosphorothioate oligonucleotide to target Navl.3 #2457.

Synthetic

WO2004016754-A2.

26-FEB-2004

14-AUG-2003; 2003WO-US025465

14-AUG-2002; 2002US-0403416P

(PHAA ) PHARMACIA CORP

Roberds SL;

WPI; 2004-203785/19.

New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

Claim 4; SEQ ID NO 2457; 417pp; English.

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound appecifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to encropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate

headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA. 8888888888

Sequence 20 BP; 1 A; 2 C; 0 G; 17 T; 0 U; 0 Other;

Gaps ö Length 20; 0.6%; Score 17; DB 1; Length 20; 100.0%; Pred. No. 1e+03; ive 0; Mismatches 0; Indels Conservative Local Similarity les 17; Conser Query Match Best Loca Matches

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2709 AAAAAAAAAAAAAAA 2725 AAAAAAAAAAAAAA 1 11 ઠ 셤

ADK74838,

BP ADK74838 standard; DNA; 20

ADK74838;

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Gaps

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0; Indels

(first entry) 20-MAY-2004

Chimeric phosphorothioate oligonucleotide to target Nav1.3 #2172

Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.

Synthetic.

WO2004016754-A2

26-FEB-2004.

14-AUG-2003; 2003WO-US025465.

14-AUG-2002; 2002US-0403416P.

(PHAA ) PHARMACIA CORP

Roberds SL;

WPI; 2004-203785/19.

New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

Claim 4; SEQ ID NO 2172; 417pp; English.

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, condition trom burns, migraine headache, cluster headache, mild-to-moderate conditions headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represente a chimeric phosphorchhoad seizure disorder with 2 MoE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target. 

Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;

Query Match

0.6%; Score 17; DB 1; Length 20;

AAQ7567

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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of
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AAQ75798 standard; DNA; 21

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                                                  Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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                                                                                                      Disclosure; Page 7; 11pp; Japanese.
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AAQ75660 standard; DNA; 21 BP.
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and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                    Reverse transcription primer used in cDNA analysis technique.
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Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                              Disclosure; Page 6; 11pp; Japanese
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                            Reverse transcription primer used in cDNA analysis technique.
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AAQ75666 standard; DNA; 21
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(first entry)
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             Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abablled reverse transcription primers (GENESEQ files AAQ75547-Q7578) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                      0.6%; Score 17; DB 1; Length 21; 100.0%; Pred. No. 1.1e+03;
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                                                                       Disclosure; Page 7; 11pp; Japanese.
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Matches 17; Conservative
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A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                      - by amplification of mRNA followed
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 9; 11pp; Japanese.
2709 AAAAAAAAAAAAAA 2725
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                                                                                               AAQ75789 standard; DNA; 21
                                                                                                                                                         (first entry)
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Matches 17; Conservative
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Gaps

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Analysis of CDNA and gene expression - by amplification of mRNA followed
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                                                                                                                                                                                                                                                                                                                               double stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                             for the analysis of cDNA comprises (a) preparing an aggregate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 0 A; 2 C; 2 G; 17 T; 0 U; 0 Other;
                                                                                                                                   (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                  by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                       Disclosure; Page 9; 11pp; Japanese.
                                                        93JP-00112515
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                                                      16-APR-1993;
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                                                                                                                                                                                                                                                                                                                 A method
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Gaps ó 0.6%; Score 17; DB 1; Length 21; 0; Indels 1.16+03; 100.0%; Pred. w. Local Similarity 100. Query Match Matches

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2709 AAAAAAAAAAAAAA 2725 17 AAAAAAAAAAAAAA 1 ઠે a

MAP3K9 marker amplification reverse primer #1057. ADW84561 standard; DNA; 21 BP (first entry) 07-APR-2005 ADW84561; RESULT 1376 ADW84561 

mixed lineage kinase; MLK; asthma; at-risk haplotype; MAP3K9; antiasthmatic; respiratory-gen.; antiinflammatory; antirheumatic; antiasthritic; antipsoriatic; neuroprotective; gastrointestinal-gen.; respiratory disease; chronic obstructive pulmonary disease; chronic betructive pulmonary disease; chronic betructive pulmonary disease;

Unidentified.

WO2005007144-A2

27-JAN-2005.

14-JUL-2004; 2004WO-US022446.

14-JUL-2003; 2003US-0487072P. 05-APR-2004; 2004US-0559611P.

(DECO-) DECODE GENETICS EHF.

Halapi E; Gurney ME, Hakonarson H,

WPI; 2005-122681/13.

Use of mixed lineage kinase family kinase inhibitor in the manufacture of a medicament for treatment of asthma associated at-risk haplotype for asthma, at-risk haplotype in MAP3K9 gene or increased MLK1 protein expression or activity.

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The invention relates to the moved use or a mixed inheade Kinase (Mikk) associated with a risk factor selected from an at-risk haplotype for associated with a risk factor selected from an at-risk haplotype for asthmatic asthmatic and the protect of a mixed in MAPIX9 gene, polymorphism in MAPIX9 nucleic acid, dysregulation of a MAPIX9 mRNA expression, dysregulation of a MAPIX9 mRNA september or identification of a mean of turber comprises a method for the diagnosis or identification of a mean of a method for the diagnosis or identification of a mean of a method for the use of a first nucleic acid molecule in a method for assessing the response to treatment with an MIX ample; a method for assessing the response to treatment with an MIX ample; a method for assessing the response to treatment with an MIX ample; a method for assessing the response to treatment with an MIX ample; and increased MIX1 brotein actid, dysregulation of in an individual with an at-risk haplotype for asthma, as above; a kit for MAPIX9 mRNA expression, increased MIX1 blochemical activity or increased transpared mixed protein isoform expression; a method for assessing the response to treatment with an MIX41 inhibitor in a target population including an individual with an at-risk haplotype for asthma, as above; a kit for assaying a sample for the presence of at least one nucleic acid capable of detecting the presence of at least one nucleic acid capable of detecting the presence of at least one mucleic acid capable of detecting the presence of at least one mucleic acid capable of detecting at least one haplotype comprising 2 or more alleles comprising; at least one mucleic acid capable of detecting at least one haplotype or assaying a sample for the presence of at least one haplotype or assaying a sample for the presence of at least one mucleic acid capable of detecting at least one mucleic acid capable of detecting at least one mucleic acid capable of detecting at least one mucleic acid approximate and reagents for detecting at least one mucle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       susceptibility to asthma. The inhibitor is also useful for the treatment of other respiratory diseases associated with MAP3K9 or other members of the JNK pathway such as chronic obstructive pulmonary disease, chronic bronchitis and other inflammatory diseases such as rheumatoid arthritis, psoriasis, multiple sclerosis and inflammatory bowel disease. This polynucleotide sequence represents a reverse primer which is used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MLK1 protein isoform expression; and in diagnosis or identification of
                                                                             The invention relates to the novel use of a mixed lineage kinase (MLK)
                          Disclosure; Fig 12; 640pp; English
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ö 0.6%; Score 17; DB 1; Length 21; 100.0%; Pred. No. 1.1e+03; ive 0; Mismatches 0; Indels Sequence 21 BP; 5 A; 7 C; 4 G; 5 T; 0 U; 0 Other; 1146 AGAGGATCATGCTGTTC 1162 17 AGAGGATCATGCTGTTC 17; Conservative Best Local Similarity Query Match Matches ઠ g

amplifying a marker of the MAP3K9 kinase, where MAP3K9 is a part of Mitogen-Activated Protein Kinase (MAPK) signal transduction pathways, of

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Gapa

AAT38295 standard; DNA; 20 RESULT 1377 AAT38295 SAKKEX

BP.

29-MAY-1997 (first entry)

AAT38295;

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PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAX36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, aslpiniquis, perihepatitis, bartholinitis, pneumogathy in breast feeding infants; and veneral lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analyzing target nucleic acid sequences, useful for population genetics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reverse phase triplex forming oligonucleotide; RP-TFO; protected nucleic acid sequence; PNAS; single nucleotide polymorphism; SNP; short tandem repeat; cancer; Factor V Leiden SNP; ss.
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Pred. No. 1.1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                           Genome sequence of Chlamydia trachomatis.
                                                                                                                                                                                                                                                    Disclosure; Page 1713; 1755pp; English.
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97FR-00015041.
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98US-0107077P.
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03-NOV-1999; 99US-0163416P.
21-DEC-1999; 99US-0171348P.
07-JUL-2000; 2000US-0216579P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 90.0
nes 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ramberg ER;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-343488/36.
                                                                                                                                                                    WPI; 1999-371125/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CYGE-) CYGENE INC. (OSTE/) OSTE C C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200132929-A1.
                                                                                     (GEST ) GENSET
                     17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-MAY-2001.
                                                                                                                              Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS05713;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting infection or contamination by Paracoccidioides - using specific primers for nucleic acid amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT38294-303 are primers specific for an unique Paracoccidioides brasiliensis sequence. The unique sequence is contained in a 110 bp fragment (AAT38293) which includes 48 nucleotides of rat beta-actin primer sequence and 62 bp of unique P. brasiliensis sequence. The unique sequence can be used as a target for detection of Paracoccidioides infection. The specific primers are used to detect P. brasiliensis infection in mammals (and to monitor treatment) or contamination of food, water, soil, manufactured products (e.g. pharmaccuticals) or biomass. The method allows early (including pre-natal) diagnosis, and since samples can be sterilised is safe to perform
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine, eye disease, conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; gential disease, pertihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; preumopathy; venereal lymphogranulomatosis; 88.
                                       primer; PCR; polymerase chain reaction; amplify; unique;
Paracoccidioides brasiliensis; rat beta-actin gene; target; detection;
monitor treatment; contamination; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
    Specific primer for unique P. brasiliensis genomic DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'Match 0.6%; Score 16.8; DB 1; Length 20; Local Similarity 90.0%; Pred. No. 1.1e+03; les 18; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer used to amplify an ORF of Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2571 GAGCTAGGAAGAGTCTACCC 2590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 16; Page 27; 39pp; English
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Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                              (UYBO-) UNIV BOSTON.
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-443205/44.
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                                                                                                                                                                                                                                                             21-MAR-1996;
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                                                                                                                                   Synthetic.
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RESULT 1378

Matches

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Gaps

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comprises hybridizing triple drug development and diagnosing cancer, comprises hyk forming oligonucleotide and probe to target sequence.

Example 2; Page 66; 141pp; English.

reverse phase triplex forming oligonucleotide, RP-TFO, (3' to the SNP) the target SNP used to analyse Factor V Leiden SNP using the method of the invention. The invention relates to analysing target nucleic acid sequences comprising restricting isolated DNA, hybridising at least one triplex forming oligonucleotide (TFO), adding a 3' to 5' exonuclease to form a protected nucleic acid sequence (FNAS) tail structure, hybridising the captured structure with a single nucleotide polymorphisms (SNP) identification probe and determining the SNP score. The methods can be sequences, to determine if they contain SNPs or short tandem repeats (STRS). The methods can be used for analysing target nucleic acid sequences, especially genomic DNA sequences, to determine if they contain SNPs or short tandem repeats (STRS). The methods can be used to detect SNPs for use in population genetics, drug development, forensics, cancer, genetic disease research, genomic analysis, diagnostics and therapeutics in humans, plants and sequence is a polypyrimidine oligonucleotide for binding a animals

Sequence 20 BP; 1 A; 1 C; 0 G; 18 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. 1.1e+03; 2; Indels 0; Mismatches 18; Conservative Query Match Best Local Similarity Matches

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2709 AAAAAAAAAAAAAAAA 2728 ઠે

20 AAATAAGAAAAAAAAAA 1 쉱

AAF83959 standard; DNA; 20 BP 06-AUG-2001 (first entry) AAF83959; RESULT 1380 CCCCCX8X4444X8X1X4X4X4X4X4X8X8X8X8X8XX

prostate; tumour; cancer; diagnostic; genetic analysis; PCTA-1; BAP28 gene fragment amplifying primer BAP28polyTcourt. PCR primer; ss. BAP28;

Homo sapiens.

WO200100669-A2.

04-JAN-2001

23-JUN-2000; 2000WO-IB001183

25-JUN-1999; 99US-0141323P. 18-JAN-2000; 2000US-0176880P.

(GEST ) GENSET,

A; Cohen-Akenine Chumakov I, ij Bougueleret Barry C,

WPI; 2001-367032/38.

New BAP28 polynucleotides and polypeptides overexpressed in prostate cancer cells for diagnosing prostate tumors, e.g. by hybridization or polymerase chain reaction assays.

Example; Page 347; 349pp; English.

The invention is directed to BAP28 polypeptides, BAP28 polynucleotide sequences and regulatory region located at the 3' and 5' ends of the BAP28 coding region. The BAP28 polypeptides can be expressed by standard recombinant methodology. BAP28 polynucleotides and polypeptides have been found to be over expressed in prostate tumour cells, therefore levels of

BAP28 expression and/or activity may be assayed (e.g. by polymerase chain reaction (PCR)) to diagnose patient suffering from or susceptible to prostate cancer. Antibodies specific for the BAP28 polypeptides are useful as diagnostic reagents. Biallelic markers of the BAP28 gene are useful in genetic analysis. Sequences AAR83934-963 represent primers for the BAP28 gene and PCTA-1 gene (the coding strand of PCTA-1 gene is on the opposite of the coding strand of BAP28) 88888888888

Sequence 20 BP; 2 A; 0 C; 1 G; 17 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. 1.1e+03; ive 0; Mismatches 2; Indels Query Match 0.6 Best Local Similarity 90.0 Matches 18; Conservative

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2703 TGTACTAAAAAAAAAAAA 2722 20 TATACAAAAAAAAAAA ð 셤

RESULT 1381

Rat protein phosphatase 2 oligo inhibitor SEQ ID No 100. ABT07486 standard; DNA; 20 BP. (first entry) 14-NOV-2002 ABT07486; ABT07486

Cytostatic; antidiabetic; antisense therapy; aberrant insulin regulation; protein phosphatase 2 catalytic beta subunit; antisense compound; cancer; hyperproliferative disorder; diabetes; inflammation; tumour; rat; ds.

Rattus norvegicus 

WO200264737-A2

22-AUG-2002

31-JAN-2002; 2002WO-US002805

09-FEB-2001; 2001US-00780045

(ISIS-) ISIS PHARM INC.

Wyatt JR; Monia BP, WPI; 2002-657588/70.

New antisense oligonucleotides targeted to nucleic acid encoding Protein Phosphatase 2 catalytic subunit beta, useful for treating diseases Phosphatase 2 catalytic subunit beta, useful for treating diseases related to Protein Phosphatase 2 catalytic subunit beta expression, such cancer aB

Example 16; Page 98; 137pp; English.

The invention relates to a novel compound 8-50 nucleotides in length targeted to a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit, where the compound specifically hybridises with an inhibite the expression of protein phosphatase 2 catalytic beta subunits, or specifically hybridises with at least an 8-nucleotide portain of an active site on a nucleic acid molecule encoding a protein phosphatase 2 catalytic beta subunit. The antisense compounds are useful for modulating the expression of protein phosphatase 2 catalytic beta subunits and for tracting diseases or conditions associated with expression of protein phosphatase 2 catalytic beta subunits, e.g. aberrant insulin regulation or diabetes or a hyperproliferative disorder, particularly cancer. The antisense compounds are also useful for diagnostics, therapeutics, prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and indistinguishing between functions of various members of a biological pathway. This polynucleotide sequence represents an oligonucleotide inhibitor of rat protein phosphatase 2 catalytic beta

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subunit mRNA levels of the invention. NOTE: This oligonucleotide contains phosphorothioate residues and has 2'- MOE wings with a deoxy gap
    88333
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Sequence 20 BP; 2 A; 10 C; 8 G; 0 T; 0 U; 0 Other;

Gaps ö 20; IndelB Length Score 16.8; DB 1; Pred. No. 1.1e+03; 0; Mismatches 405 CCGCGGCCGCCGCCCG 424 0.6%; Query Match
Best Local Similarity 90.0
Matches 18; Conservative 셤 ઠે

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1 caecegecaecececeece 20

ABZ85669 standard; DNA; 20 BP (first entry) 17-0CT-2003 ABZ85669; RESULT 1382 

Human oligonucleotide sequence.

antinflammatory steroid, ubiquinone, antinflammatory; antiallergic, antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antiense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds. antisense; lung dysfunction; nasal airway dysfunction; Human;

Homo sapiens.

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar Pabalan J, Katz E, i Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Li Y, Nyce JW, I Miller S,

ä

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.

Claim 15; SEQ ID NO 911; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' or and 3' intron-exon junctions, 5' or 3' end genomic flanking regions, 5' or and 3' intron-exon junctions of zegions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antishammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject, stissue, or treating bronchoconstriction, lung ainflammatton, lung allergiae, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed

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specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                       Gaps
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0
                                                                          Length 20;
                                                                                                     IndelB
                                             Sequence 20 BP; 0 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
                                                                     Score 16.8; DB 1;
Pred. No. 1.1e+03;
0; Mismatches 2;
                                                                          0.6%;
                                                                                                        18; Conservative
                                                                                         Local Similarity
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2709 AAAAAAAAAAAAAAAA 2728 20 ð 셤

1383 ABZ89176

ABZ89178 standard; DNA; 20 BP

ABZ89178;

17-OCT-2003 (first entry)

Human oligonucleotide sequence

Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens.

WO200285308-A2

31-OCT-2002.

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P.

(EPIG-) EPIGENESIS PHARM INC.

Aguilar D; Pabalan J, Katz E, Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; Miller S, Ř Nyce

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone 

Disclosure; SEQ ID NO 4420; 872pp; English.

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, c' initiation codon, coding regions, 5' or 3' end genomic flanking regions, c' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal alrway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiathmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, indo or mallignant disease or condition, also antiinflammatory steroid in a subject, for reducing or depleting levels of for enhancing the prophylactic or therapeutic respiratory effect of an antismanatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of demosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject is tissue, or a respiratory disease or condition. Unug allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed

Gaps

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end ganomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory antiallergic, antiasthmatic, hypotensive, fumunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of denosine receptor, producing sensitivity to adenosine, reducing levels of the printed repetence or lung surfactent in a subject, increasing levels of biquinone or lung surfactent in a subject or respiratory disease or condition.

Column surfactent in a subject or respiratory disease or condition. Ung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating ailments associated with impaired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiathautory steroid; ubiquinone; antiathautory; antiallergic; antiathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; admissense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
  was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense; lung dysfunction; nasal airway dysfunction;
                                                                                                   Length 20;
                                                                                                                                             Indels
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                                                           Sequence 20 BP; 13 A; 0 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                         2,
                                                                                                 Score 16.8; DB 1;
Pred. No. 1.1e+03;
              at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; SEQ ID NO 777; 872pp; English.
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, Shahabuddin S;
                                                                                                                                                                                 2702 TTGTACTAAAAAAAAAA 2721
                                                                                                                                                                                                                          TTGTTTTAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human oligonucleotide sequence.
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                                                                                                   0.68;
                                                                                                                      90.06
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                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                             Conservative
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                                                                                                                    Local Similarity
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Tang
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                   ABZ85535;
                                                                                                 Query Match
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                                                                                                                                       Matches
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comparising oligonucleotides, effective for alleviating contribution describes and bronchoconstriction, respiratory tract inflammation allergies and bronchoconstriction, respiratory tract inflammation allergies and bronchoconstriction or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating capression of a target polypeptide associated with lung alrawy or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antisathmatic, instructions density, immunosuppressive and cytostatic activity, is a halfage of the composition is useful for preventing or treating a respiratory, lung or mailgnant disease. The administered composition composition may administered to reduce the production
                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant deptetion; antiallergic; antialfammatory; antialantacic; analysesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention describes a novel composition (a) a first active agent
                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide containing less percentage of adenosine, targeted t
nucleic acids associated with lung airway or lung dysfunction, and
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy; hyposecretion;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense
                                                                                       Score 16.8; DB 1; Length 20;
Pred. No. 1.1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pabalan J,
                                                    Sequence 20 BP; 18 A; 0 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                  AI122807-derived oligonucleotide SEQ ID 4420.
                   at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; SEQ ID NO 4420; 763pp; English.
                                                                                                                                                                   2709 AAAAAAAAAAAAAAAAA 2728
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                                                                                                                                                                                        Sandrasagra A,
                                                                                                                                                                                                                                                                                                   BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC.
                                                                                           0.68;
                                                                                                          90.06;
                                                                                                                                                                                                                                                                                                   ABD25408 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                              18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bronchodilating agent.
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Tang L,
                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Miller S,
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                                                                                           Query Match
                                                                                                              Local
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                                                                                                                                                                                                                                                              RESULT 1385
                                                                                                                              Matches
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or availability, or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhintis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted uncleic acids associated with lung airway or lung dysfunction, and
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                                                                                                                                                                                                                                                                                                                                               0.6%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                       BP; 13 A; 0 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human stanniocalcin-derived oligo SEQ ID 777.
                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Li Y, Sandrasagra A, Ka
Tang L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                              2702 TIGIACTAAAAAAAAAA 2721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 TrGTTTTAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABD21765 standard; DNA; 20 BP
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hes 18; Conservative
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                                                                                                                                                                                                                                                                                                         Sequence 20
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABD21765;
                                                                                                                                                                                                                                                                                                                                                   Query Match
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congrated depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

The invention also describes a kit, that comprises: (a) a delivery containers, (b) the oligonucleotides, (c) evice, in separate containers, (b) the oligonucleotides, (c) enstructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition confiction, and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung confiction and separate or conditions and/or bronchoconstriction, and separate or conditions and as a such as pulmonary disease, pulmonary confictions, palm, cystic fibrosis, allergic rhinitis, pulmonary configuration rejection, pulmonary infections, bronchiis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system environment and thereby, to prevent any unwanted effects due to it ö Human, antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; anniallergic, antiinflammatory; antisathmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic fibritis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer. Gaps ö ; Aguilar Length 20; 2; Indels Katz E, Pabalan J, Sequence 20 BP; 18 A; 0 C; 2 G; 0 T; 0 U; 0 Other; Score 16.8; DB 1; Pred. No. 1.1e+03; Human stanniocalcin-derived oligo SEQ ID 911. 0; Mismatches Li Y, Sandrasagra A, Ka Tang L, Shahabuddin S; 2709 AAAAAAAAAAAAAAAA 2728 1 AAAAAAAGAAAAAAA 20 BP. 23-APR-2002; 2002WO-US013143. 24-APR-2001; 2001US-0286036P. (EPIG-) EPIGENESIS PHARM INC 0.6%; 90.06; ABD21899 standard; DNA; 20 29-JUL-2004 (first entry) Query Match
Best Local Similarity 90.09 WPI; 2003-093058/08 WO200285309-A2. Homo sapiens. 31-OCT-2002. Miller S, ABD21899; Nyce JW, RESULT 1387 음 ઠે

Homo sapiens.

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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating by threation, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that compositions of the invention also describes a kit, that composition of the invention as antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary viseconstriction, and seems of inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary disease, pulmonary contained adenosine content of the anti-sense oligos corresponding to the inflammation rejection, pulmonary infections, brank, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                      Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                             invention describes a novel composition (a)
                                                                                                                                                      Claim 15; SEQ ID NO 911; 763pp; English
                                                                                                      bronchodilating agent.
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Sequence 20 BP; 0 A; 2 C; 0 G; 18 T; 0 U; 0 Other;

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Score 16.8; DB 1; Length 20;
Pred. No. 1.1e+03;
0; Mismatches 2; Indels
                                                         2709 AAAAAAAAAAAAAAAAA 2728
                             ;
0
  0.6%;
                Best Local Similarity 90.0
Matches 18; Conservative
  Query Match
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Gaps

## 20 AGAAAAAAAGAAAAAAA 1 셤

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ADH70655 standard; DNA; 20 BP.
                  25-MAR-2004 (first entry)
            ADH70655;
1388
   ADH70655
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Human Vbeta gene repeat sequence #445.

human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; addison's disease; atrophic gastritis; disease; repelse atrophic gastritis; Alzheimer, s disease; hypersensitivity disease; type I hypersensitivity disease; type I hypersensitivity; doodpasture's syndrome; type IV hypersensitivity; doodpasture's syndrome; type IV hypersensitivity; doodpasture's syndrome; HIV; fungal infection; dandda; parasitic infection; enfistesome; filaria; bacterial infection; Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;

breast cancer; ds

Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a 94US-00309335. 95US-00531241. 99US-00263959 WPI; 2004-059052/06. Hood LE, Rowen L; (HOOD/) HOOD L E. (ROWE/) ROWEN L. US2002150891-A1. 05-MAR-1999; 19-SEP-1994; 19-SEP-1995; 17-OCT-2002. Vbeta gene 

a first active agent,

The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each 'Detag gene, 'Detagn's or converse again, 'Detagn's or and diagnosing and treating T-cell associated diseases concluming autoimmune diseases, 'degenerative nervous system diseases, 'Grarophing autoimmune diseases, 'Hypersensitivity diseases, infectious disease, and neoplastic disease. Autoimmune diseases include Addison's disease, arrophic gastritis. Degenerative nervous system diseases include multiple carrophic gastritis. Degenerative nervous system diseases include multiple callergies, Type II hypersensitivities such as those present in Coodpasture's syndrome and Type IV hypersensitivities such as those caused by the yeast genus Candid, parasitic infections such as those caused by the yeast genus Candid, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by shitted and bacterial infections such as those caused by who batterium hopping and hyphoproliferative diseases such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence. Disclosure; SEQ ID NO 849; 164pp; English. lymphomas and cancers such as leukaemias,

ö Gaps .. 0 0.6%; Score 16.8; DB 1; Length 20; 0.0%; Pred. No. 1.1e+03; ve 0; Mismatches 2; Indels Sequence 20 BP; 18 A; 0 C; 2 G; 0 T; 0 U; 0 Other; 2709 AAAAAAAAAAAAAAAA 2728 90.06; Query Match
Best Local Similarity 90.0
Matches 18; Conservative 8

1 AAAAGAAAAAAAAAAAAA 20 RESULT 1389 셤

ADL01298 standard; DNA; 20 BP

ADL01298

ADL01298;

vascular endothelial growth factor; ss; antisense compound; phosphorochioate linkage; 2'-O-methoxyethyl sugar moiety; 5-methylcytosine; antisense oligonucleotide; diabetes; immunological disorder; cardiovascular disorder; neurological disorder; ischaemia; reperfusion injury; cancer; angiogenic disorder; haemangioma; Human VEGF co-regulated chemokine-1 DNA antisense oligonucleotide #831. Human; VEGF co-regulated chemokine-1; VCC-1; 20-MAY-2004 (first entry) 

01-JUL-2004 (first entry)

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The invention relates to an antisense compound targeted to a nucleic acid molecule encoding human vascular endothelial growth factor (VEGF) co-
cegniated chemokine-1 (VCC-1), and which specifically hybridises with and inhibits the expression of VCC-1. The invention also relates to a composition comprising the antisense compound, a method of inhibiting the expression of VCC-1 in cells or tissues compound, a method of inhibiting the calls or tissues with the antisense compound and a method of treating a human having a disease or condition associated with VCC-1 comprising or tissues with the antisense compound and a method of treating a human comprise at least one modified comprises at least one modified comprises at least one modified comprises at least one modified sugar moiety, preferably a 2'-0-
comprises at least one modified sugar moiety, preferably a 2'-0-
methoxyethyl sugar moiety, and at least one modified nucleobase,
comprises at least one condition associated with VCC-1, such as diabetes,
a chimeric oligonucleotide. The antisense oligonucleotide preferably is a chimeric oligonucleotide. The antisense oligonucleotide for condition associated with VCC-1, such as diabetes,
an immunological disorder, a cardiovascular disorder, a neurological
disorder, ischaemia, reperfusion injury, cancer or an angiogenic
confusorder, e-g. haemangiona, tumour angiogenseis, rheumatoid arthritis,
atherosclerosis, psoriasis or fibrosis after myocardial infarction. VCC-1
antisense oligonucleotides may also be used for wound healing, for treatment of this sequence represents an antisense
corresponder, e-g. humour angiodenseis, rheumangion, for treatment of lung or liver fibrosis or for cregeneration, for treatment of lung or liver fibrosis or for management of atrial fibrillarion. This sequence represents an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide targeted to DNA encoding the human VCC-1 polypeptide of
tumour angiogenesis; rheumatoid arthritis; atherosclerosis; psoriasis; fibrosis; myocardial infarction; wound healing; bone fracture; cartilage damage; tissue regeneration; organ regeneration; periodontal disease; gut regeneration; atrial fibrillation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compounds targeted to a nucleic acid molecule encoding vascular endothelial growth factor co-regulated chemokine-1 (VCC-1), useful for treating VCC-1-associated disorders, e.g. diabetes or a neurologic disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 831; 336pp; English.
                                                                                                                                                                                                                                                                                             19-AUG-2003; 2003WO-US025891.
                                                                                                                                                                                                                                                                                                                                                19-AUG-2002; 2002US-0404484P.
                                                                                                                                                                                                                                                                                                                                                                                                PHAA ) PHARMACIA CORP.
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                                                                                                                                                                                                                                             26-FEB-2004
                                                                                                                                          Homo
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0.6%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. 1.1e+03; ive 0; Mismatches 2; Indels
                                                                  GTGAATTTTAAAAATATTTA 963
                                                                                               Greacatriaaaaatatria 20
Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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ADM14429 standard; DNA; 20
                                                           ADM14429;
RESULT 1390
                ADM14429
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                          /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:616.
                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                  'mod_base= OTHER
'note= "2'-0-methoxyethyls"
                                                                                                                                                                Location/Qualifiers
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modified_base
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                                                                                                                                    Homo sapiens
                                                                                                                                             Synthetic
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New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g inflammation, Alzheimer's disease, arthritis, diabetes, cancer or ischemia.

25-SEP-2003; 2003WO-US030374 25-SEP-2002; 2002US-0413549P.

WO2004028458-A2

08-APR-2004.

(PHAA ) PHARMACIA CORP

WPI; 2004-305094/28.

Gierse JK;

Claim 4; SEQ ID NO 616; 132pp; English.

antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiarmatory, neuroprotective, nootropic, antidarthitic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9d34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibite its expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MPGES-1 chimeric antisense oligonucleotides and anishense compounds have cytostatic, ophthalmic, immunological, cardiovascular or neurological disorder.

Sequence 20 BP; 1 A; 1 C; 2 G; 16 T; 0 U; 0 Other;

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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA sample containing at least one segment of (1); and cast exemining the length of the resulting amplicon(8). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or concerned pML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a diseases and for prediagnosis of such diseases. especially prion diseases concerned diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperhermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, comparation for prediagnosis of such diseases that encode milk proteins, particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion comparising a polymorphic microsatellite locus.
                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                 gene typing; polymorphic microsatellite loci; PML; disease; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow; microsatellite; PCR; primer; ss.
                                             ö
   Length 20;
   Score 16.8; DB 1; Length 2
Pred. No. 1.1e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                               prion protein microsatellite locus primer #70.
                                         0; Mismatches
                                                                             2705 TACTAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 28; 64pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Han Y;
                                                                                                                 20 TGCCAAAAAAAAAAAAA 1
                                                                                                                                                                                                                ADO81058 standard; DNA; 20 BP
0.64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                        29-JUL-2004 (first entry)
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Best Local Similarity 90.0°
Matches 18; Conservative
                                         18; Conservative
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                       Local Similarity
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   Query Match
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The present invention relates to a novel pharmaceutical composition comprising: (a) an agent capable of modulating an expression level or protein activity of a cancer-related transmembrane protein (CRTP) or gene; an antibody specific for a CRTP, or a T cell activated by a CRTP; and C(b) a carrier. The pharmaceutical composition may also comprise a polynucleotide capable of inhibiting or decreasing the expression of the CRTP practice or an antisense mechanism. The CRTPs of the invention are selected from ABCC4, CZOORTION, CCRONID, CDHG, CST, ENPP3, CT-11856, GPR54, HAVCR1, SLC6A3, SLC30A4, TRG, and TRPM4. The CRTPs of the pharmaceutical composition is useful for treating cancer, e.g. colon cancer, lung cancer, neast cancer, liver cancer, kidney cancer, stomach cancer, and sephageal cancer. The present sequence is a target oligonucleotide from one such CRTP for which short interfering cancer, stomach directly from who are produced. Note: The sequence data for this patent did cont form part of the printed specification, but was obtained in electronic format directly from WIDD at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition comprising an agent capable of modulating an expression level or protein activity of a gene, e.g. ABCC4, or a T cell activated by the polypeptide or antibody, and a carrier, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                 Cytostatic; Gene therapy; Vaccine; RNA Interference; cancer; ss.
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                                                                                                                                                                                                                  TRPM4 target oligonucleotide, SEQ ID 14539
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Howes SH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2121 ACCTGGAGGCCTTGGCTTG 2140
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ВР
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                                                                                                         ACL53467 standard; DNA; 21
                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                        Homo sapiens.
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Gaps

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0.6%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. 1.1e+03; tive 0; Mismatches 2; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises antisense oligonucleotides that are targeted to nucleic acid molecules encoding human signal transducers and activators of transcription 3 (STAT3). The antisense oligonucleotides of the invention inhibit expression of human STAT3. The antisense of the oligonucleotides of the invention are useful for treating and preventing inflammatory diseases (e.g. rheumatoid arthritis) and cancers (e.g. and lymphomas). The present DNA sequence represents a human STAT3-specific antisense oligonucleotide. NOTE: The present sequence has a phosphorothioate backbone, 2'-MOE wings and a deoxy gap.
                                                                                                                                                                                                                                                                                                                                  New antisense compound, useful for treating or preventing inflammatory diseases (e.g. rheumatoid arthritis) and cancers (breast, prostate, head and neck, and brain cancer, myelomas, melanomas, leukemias, and
                                                    antisense oligonucleotide; antisense therapy; inflammation; antinflammatory; rheumatod arthritis; antiarthritic; antirheumatic; cancer; ytostatic; breast tumor; prostate tumor; head & neck tumor; brain tumor; multiple myeloma; melanoma; leukemia; lymphoma; STAT3; ss; phosphorothioate; 2'-O-methoxyethyl; 2'-MOE wing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human tumour necrosis factor receptor mRNA; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligomer TNFR941 for forming triplex with HUMNFR target duplex.
                                Human STAT3-specific antisense oligonucleotide - SEQ ID 401.
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                                                                                                                                                                                                                                                                                                                                                                                           Example 22; SEQ ID NO 401; 149pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2438 AAGAAGCAGGAGCTGCTGGA 2457
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14-NOV-2003; 2003US-00713139.
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                                                                                                                            Homo sapiens
         16-JUN-2005
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07-DEC-1992
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The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor receptor mRNA beginning at nucleotide 2354 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets. e.g. HPV, HER, HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PD field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.
                                                                                                                                                                   /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Froehler B, Krawczyk S, Matteucci MD, Milligan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 18 BP; 1 A; 1 C; 0 G; 16 T; 0 U; 0 Other;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 12; Page 72; 77pp; English.
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                                                                           mod_base= m5c
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
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                                                                                                            modified base
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17-APR-1991;
17-APR-1991;
27-SEP-1991;
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08-APR-1991;
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                                                                                                                                                                                                                 Producing binary sequence tags, useful for analyzing nucleic acid sequence tags, gene expression or gene-expression patterns, involves generating nucleic acid fragments, which are mixed with offset adaptors
                                                                                                                                                                                                                                                                                                         The present invention describes a method of producing binary sequence
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                                                                                                                                                                    Latimer DR;
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Pred. No. 1.1e+03;
                                                                                                                                                                  Feng L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Feng L,
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                                                                                                                                                                  Lizardi PM,
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                                                                          11-AUG-2000; 2000WO-US022164.
                                                                                                    13-AUG-1999; 99US-0148870P.
06-APR-2000; 2000US-00544713.
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2000US-00544713
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                                                                                                                                                                  Roth ME,
                                                                                                                                                                                                                                                        and adaptor-indexers.
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                                                                                                                                                                                         WPI; 2001-202878/20.
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                                                                                                                                       (UYYA ) UNIV YALE.
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                          WO200112855-A2.
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06-APR-2000;
                                                                                                                                                                 Kaufman JC,
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                                                22-FEB-2001
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 Synthetic.
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                                                                                                                           The present invention describes a method of producing binary sequence tags from nucleic acid fragments in a sample, involving incubating the sample with cleaving reagents, mixing offest adaptors with the sample, incubating with more cleaving reagents and mixing the sample with adaptor -indexers where the adaptors are coupled to binary sequence tags. The method is useful in sequence analysis, including analysis and comparison of gene expression, nucleic acid samples and genomes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for providing a profile of mRNA molecules present in a sample. The method comprises generating two independent patterns characteristic of the population of mRNA molecules expressed in the sample and analysing the patterns using a combinatorial algorithm, comparing gene expression by different or same cell types under different conditions, and identifying genes having a role in various cellular processes. The method is useful for the analysis and identification of transcribed genes, and fingerprinting. The method can be used to identify genes which play a role in determining various
                   sequence tags, gene expression or gene-expression patterns, involves generating nucleic acid fragments, which are mixed with offset adaptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Providing mRNA profile, by generating two independent patterns
characteristic of sample mRNA population, analyzing patterns, comparing
gene expression by cell types under varied conditions, and identifying
                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identification of transcribed gene; mRNA profile, gene expression; cellular process; fingerprinting; susceptibility to external factor; development; disease; PCR; primer; ss.
Producing binary sequence tags, useful for analyzing nucleic acid
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                                                                                                                                                                                                                                                                                                                      Score 16.4; DB 1; Length 18; Pred. No. 1.1e+03;
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                                                                                                                                                                                                                                                                                   Sequence 18 BP; 0 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                          Disclosure; Page 100; 101pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                 2706 ACTAAAAAAAAAAA 2723
                                                                                                                                                                                                                                                                                                                                                                                                                                      18 ACAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK13935 standard; DNA; 18 BP.
                                                                                                                                                                                                                                                                                                                    0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUL-2000; 2000GB-00018016.
21-JUL-2000; 2000US-0219925P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-2001; 2001WO-IB001539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GLOB-) GLOBAL GENOMICS AB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ernfors P,
                                                                                                                                                                                                                                                                                                                                                           17; Conservative
                                                       and adaptor-indexers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-217065/27.
                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200208461-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Linnarsson S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK13935;
                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genes.
                                                                                                                                                                                                                                                                                                                                                           Matches
FFF7X8XX000000X8
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The invention relates to determining the presence of and/or identifying a present in a sample. The method involves assigning to gene fragments gene candidates within a database by comparing signals in the database with the database comprising data representing mRNAs with known colly sites and/or 'virtual genes' representing a possible polya sites and/or 'virtual genes' representing a possible polyadenylation site within an actual gene. The method is useful for determining the presence of and/or identifying a polyadenylation site or alternative polyadenylation sites within a sequence of a transcribed gene or sequences of transcribed gene variants present or potentially present in a sample, in identifying gene variants present or potentially present in a sample, in identifying gene variants that occur in a population of nucleic acid molecules, especially in identifying or discovering polya internative polya site usage or determining poly site usage or determining poly site usage or determining poly site usage in a nucleic acid sample, and gene variants arising from alternative polya sites. The present sequence
                                                                                                                                                                                                  ö
cellular processes, including susceptibility to external factors, development, and disease. The present sequence for a PCR primer is used in the production of a single pattern characteristic of a sample, employing a Type II restriction enzyme (i.e. HaeII) in the methods of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present in a sample comprises assigning to gene fragments gene candidates within a database by comparing signals in the dataset with the database.
                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining polyadenylation sites within transcribed gene sequences
                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence of a double stranded product DNA fragment.
                                                                                                                                                          Score 16.4; DB 1; Length 18;
Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gene variant identification; restriction enzyme; HaeII; ds.
                                                                                                                                                                                                1: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ernfors P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 18 BP; 0 A; 1 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                     Sequence 18 BP; 0 A; 1 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Linnarsson S,
                                                                                                                                                                                                                                       2707 CTAAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example; Fig 2; 81pp; English.
                                                                                                                                                                                                                                                                            CGAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                                                                          0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JAN-2003; 2003WO-IB000255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2002; 2002US-0352245P
                                                                                                                                                                                                                                                                                                                                                                             ACF36339 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GLOB-) GLOBAL GENOMICS AB.
                                                                                                                                                Query Match
Best Local Similarity 94.44
...-hes 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oldin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-627619/59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                    ACF36339;
                                                                                                                                                                                                                                                                                                                                        RESULT 1398
                                                                                                                                                                                                                                                                                                                                                             ACF36339,
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Producing a population of double-stranded product DNA molecules, useful for mRNA profiling, comprises amplification by nested polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to producing a population of double-stranded product DNA molecules comprising amplification by a nested PCR method. The method is useful in profiling mRNA transcribed in a system under investigation. The oligonucleotides are used as size standards in electrophoresis, and as internal controls allowing for calculation of relative amounts of material present. The present sequence represents adouble stranded product DNA, which aids in outlining an approach to production of a single pattern characteristic of a sample, employing a type II restriction enzyme (HaeII)
                                                                                                                                                                                                               Nucleic acid manipulation; mRNA profiling; polymerase chain reaction; electrophoresis; type II restriction enzyme; HaeII; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mitogen activated protein kinase siNA oligonucleotide SEQ ID NO:163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.6%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Pihlak A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                      Nucleotide sequence of a double stranded product DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 18 BP; 0 A; 1 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Metsis A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     Bauren G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2707 CTAAAAAAAAAAAAA 2724
2707 CTAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Fig 1; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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                           18 CGAAAAAAAAAAAAA 1
                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                    28-JAN-2003; 2003WO-IB000843.
                                                                                                                                                                                                                                                                                                                                                                             29-JAN-2002; 2002US-0352215P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.48;
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                                                                                        364/c
ACF36364 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE29541 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                        (GLOB-) GLOBAL GENOMICS AB.
                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       Ernfors P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-618365/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                         WO2003064691-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                     Linnarsson S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2004
                                                                                                                                                           04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Montelius A;
                                                                                                                                                                                                                                                                                                                      07-AUG-2003.
                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                               ACF36364;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                       ACF36364,
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Gaps

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0.6%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.1e+03; vative 0; Mismatches 1; Indels

Best Local Similarity 94.4 Matches 17; Conservative

Query Match

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that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, itssue explants or organisms by introduction of MAPK genes in cells, itssue explants or organisms by introduction of SINA; (2) kits for in vitro or in vivo delivery of SINA; (3) conjugates and/or complexes of SINA; and (4) vectors that express SINA and cells containing these vectors. MAPK SINAs have cytostatic, anorectic, antidabetic, antiminamatory, antiathmatic, immunosuppressive, antibacterial, antirheumatic, antipachatic and gastrointestinal activities. The MAPK SINAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I complete the activities, and inflammatory diabetes types in cells, theumatoid arthritis, psoriasis and inflammatory bowel disease). They can also be used for drug screening; pharmacogenomics; containing intention entities that the containing subarmacogenomics; containing containing that the containing special containing cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          represents a MAPK sinA which is used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a short interfering nucleic acid (siNA)
short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidabetic; antiinflammatory; antiasthmatic; immunosuppressive; antibacterial; antirheumatic; antiarthritic; antipacratic; gaetrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; rheumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated protein kinase genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK siNA which in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chowrira B;
                                                                                                                                                                                                               genetic engineering; pharmacogenomic; gene mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haeberli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 163; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beigelman L, Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JAN-2003; 2003WO-US002510.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-689980/65.
                                                                                                                                                                                                                                                                                                                                     WO2003072590-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-2002;
06-JUN-2002;
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05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-SEP-2002;
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                                                                                                                                                                                                                                                                             Synthetic.
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ö Gaps . 0 0.6%; Score 16.4; DB 1; Length 19; 18.9%; Pred. No. 1.1e+03; 1; Indels Sequence 19 BP; 16 A; 1 C; 0 G; 0 T; 2 U; 0 Other; 1; Mismatches 88.98; Query Match Best Local Similarity 88.9% ....hes 16; Conservative

2708 TAAAAAAAAAAAAAA 2725 CAAAAAAAAAAAAAAA 19

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ADE29704 standard; RNA; 19 ADE29704; ADE29704/ ID ADE2 XX AC ADE2

RESULT 1401

(first entry) 29-JAN-2004

Mitogen activated protein kinase sinA oligonucleotide SEQ ID NO:326

short interfering nucleic acid; siNA; downregulation; inhibition; mitogen-activated protein kinase; MAP kinase; MAPK; RNA interference; cytostatic; anorectic; antidiabetic; antinflammatory; antiasthmatic; immunosuppressive; antibacteria; antirheumatic; antiarthritic; antipsoriatic; gastrointestinal; obesity; diabetes; tumour; inflammatory disease; asthma; septic shock; heumatoid arthritis; psoriasis; inflammatory bowel disease; drug screening; psoriasis; inflammatory bharmacogenomic; gene mapping; ss.

Synthetic.

WO2003072590-A1.

04-SEP-2003.

28-JAN-2003; 2003WO-US002510

20-FEB-2002;

11-MAR-2002, 2002US-0363124P.
06-UUN-2002, 2002US-03867B2P.
29-AUG-2002, 2002US-04067B4P.
05-SEP-2002, 2002US-0409293P.

(SIRN-) SIRNA THERAPEUTICS INC.

15-JAN-2003; 2003US-0440129P

Chowrira B; Haeberli P, Mcswiggen J, Beigelman L, Usman N,

WPI; 2003-689980/65.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of mitogen-activated diagnosis of cancer, or protein kinase genes.

Example 3; SEQ ID NO 326; 164pp; English.

that downregulates expression of a mitogen-activated protein kinase (MAPK) genes by RNA interference. Also described: (1) a method for modulating expression of MAPK genes in cells, tissue explants or organisms by introduction of siNA; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; and (4) vectors that express siNA and cells containing these vectors. MAPK siNAs have cytostatic, anorectic, antibacterial, antihalammatory, antiasthmatic, immunosuppressive, antibacterial, antihalammatory, antiasthratic, in antipaciatic and gastrointestinal activities. The MAPK siNAs can be used to modulate the expression of MAPK genes, in cells, tissue explants or organisms, e.g. for treating obesity; diabetes types I and II; a wide range of tumours, and inflammatory diseases (asthma, disease). They can also be used for drug screening; diagnosis; target disease). They can also be used for drug screening; pharmacogenomics; check, theumatoria shorts genetic engineering; pharmacogenomics; chicklication and validation; genetic engineering; pharmacogenomics; studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents a MAPK siNA which is used in the exemplification of the present invention. The present invention describes a short interfering nucleic acid (siNA) 

Sequence 19 BP; 2 A; 0 C; 1 G; 0 T; 16 U; 0 Other;

Gaps ö 0.6%; Score 16.4; DB 1; Length 19; 14.4%; Pred. No. 1.1e+03; ve 0; Mismatches 1; Indels 94.48; 17; Conservative Local Similarity Query Match Matches

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2725 2708 TAAAAAAAAAAAAAA

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27-JAN-2005

ADU64845

RESULT 1402

11-NOV-2004

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The invention relates to a chemically synthesized double stranded short interfering nucleic acid (siNA) molecule (I) that directs cleavage of a c-JUN RNA through RNA interference (RNA), where one strand of the siNA molecule comprises nucleotide sequence having sufficient complementarity to the c-JUN RNA for the siNA molecule to direct cleavage of the c-JUN RNA through RNA interference. (I) is useful for inhibiting mitogen activated protein kinase gene (e.g., c-JUN, JNK1, JNK2, JNS, BRK1 or BKK2) expression associated with diseases e.g., inflammatory disease, autoimmune disease, allorgy, cancer. (I) exhibits improved RNA interference activity and nuclease resistance. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mitogen activated protein kinase gene expression e.g., c-JUN associated with diseases e.g., inflammatory disease or autoimmune disease.
                                                           RNA interference; mitogen activated protein kinase inhibitor; inflammation; immunosuppressive; immune disorder; autoimmune disease; allergy; antiallergic; cytostatic; neoplasm; cancer; ss; siRNA; gene silencing; small interfering RNA; MAP kinase inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88; PCR; Cardiant; Dermatological; Gastrointestinal; Hemostatic; Respiratory-Gen; Nootropic; Neuroprotective; Uropathic; Cytostatic; Antiinflammatory; AdipoR1-inhibitor; AdipoR1-activator; G protein coupled receptor; AdipoR1; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel short interfering nucleic acid molecule useful for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.4; DB 1; Length 19;
Pred. No. 1.1e+03;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 16 A; 1 C; 0 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Usman N, Haeberli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              represents a human MAP kinase 1/ ERK2 siRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 163; 322pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human AdipoR1 reverse primer, primer2.
                   Human MAP kinase 1/ ERK2 siRNA #163.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SIRN-) SIRNA THERAPEUTICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                        30-APR-2003; 2003US-00427160.
23-MAY-2003; 2003US-00444853.
23-OCT-2003; 2003US-0053059.
24-NOV-2003; 2003US-00720448.
                                                                                                                                                                                                                                                                                                                     23-APR-2004; 2004WO-US012517.
                                                                                                                                                                                                                                                                                                                                                                   2003US-00424339.
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88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JAN-2004; 2004US-00757803
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JUN-2005 (first entry)
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Best Local Similarity 88.99
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                                                                                                                                                                                                                             WO2004097020-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J,
                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                   25-APR-2003;
                                                                                                                                                                                                                                                                        11-NOV-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polisky B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADZ00541;
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a chemically synthesized double stranded short interfering nucleic acid (siNA) molecule (1) that directe cleavage of a c-JUN RNA through RNA interference (RNA), where one strand of the siNA molecule comprises nucleotide sequence having sufficient complementarity to the c-JUN RNA for the siNA molecule to direct cleavage of the c-JUN RNA through RNA interference. (1) is useful for inhibiting mitogen activated protein kinase gene (e.g., c-JUN, JNK1, JNK2, p38, ERK1 or succious associated with diseases e.g., inflammacrory disease, autoimmune disease, allergy, cancer. (1) exhibits improved RNA interference activity and nuclease resistance. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel short interfering nucleic acid molecule useful for inhibiting mitogen activated protein kinase gene expression e.g., c-JUN associated with diseases e.g., inflammatory disease or autoimmune disease.
                                                                                                                                                                                                                           RNA interference; mitogen activated protein kinase inhibitor; inflammation; immunosuppressive; immune disorder; autoimmune disease; allergy; antiallergic; cytostatic; neoplasm; cancer; ss; siRNA; gene silencing; small interfering RNA; MAP kinase inhibitor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J, Beigelman L, Usman N, Haeberli P, Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.6%; Score 16.4; DB 1; Length 19;
94.4%; Pred. No. 1.1e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 19 BP; 2 A; 0 C; 1 G; 0 T; 16 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           represents a human MAP kinase 1/ ERK2 siRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 326; 322pp; English.
                                                                                                                                                                                     Human MAP kinase 1/ ERK2 siRNA #326.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2708 TAAAAAAAAAAAAAA 2725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2003US-00427160.
2003US-00444853.
2003US-00693059.
2003US-00720448.
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                                               ADU64845 standard; RNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best_Local Similarity 94.4%;
Matches 17; Conservative
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                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-MAY-2003;
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24-NOV-2003;
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Polisky B;

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Chowrira

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Gaps

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27-JAN-2005

ADU64682;

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ઠે g RESULT 1403

ADU64682

Query Match

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This sequence represents a primer which was used for relative quantitation of the distribution of the G protein coupled receptor, AdipoR1, mRNA in cells and tissues. The method of the invention for dermatological, therapeutic agents useful for treating cardiovascular, dermatological, gastroenterological, hematological, respiratory, neurological or urological diseases, cancer or inflammation in a mammal comprises contacting a test compound with an AdipoR1 polypeptide and detecting binding of the test compound to the AdipoR1 polypeptide and detecting binding of the test compound to the AdipoR1 polypeptide and further method is included for diagnosing any of the diseases cited above in a mammal comprising determining the amount of an AdipoR1 polypucleotide in a sample taken from the mammal and determining the amount of AdipoR1 polynucleotide in healthy and/or diseased mammals. Regulators of AdipoR1 activity are useful for regulating AdipoR1 activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                        Screening for therapeutic agents useful for treating cardiovascular, dermatological, respiratory or neurological diseases, cancer or inflammation in a mammal comprises contacting a test compound with a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              spinal cord injury; short interfering RNA; siRNA; RNA interference; gene silencing; RNA cleavage; Fas; vulnerary; ds.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.6%; Score 16.4; DB 1; Length 19; 94.4%; Pred. No. 1.1e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 1 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; SEQ ID NO 4; 135pp; English.
                                                                                                                                                                                                                                             Geerts A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in a mammal having such diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1895 GTGCCACAGGAAGAG 1912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AEA99304 standard; RNA; 19 BP
                                                                                                                                                                      27-SEP-2003; 2003EP-00021897,
                                                                                                                                    16-SEP-2004; 2004WO-EP010384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-JUN-2004; 2004US-00871222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAY-2001; 2001US-0292217P.
                                                                                                                                                                                                    (FARB ) BAYER HEALTHCARE AG.
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                                                                                                                                                                                                                                          Brueggemeier U,
                                                                                                                                                                                                                                                                                                                                                                AdipoR1 polypeptide.
                                                                                                                                                                                                                                                                          WPI; 2005-254243/26.
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Matches 17; Conserv
                                                                  WO2005031346-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2005119212-A1.
                   Homo sapiens.
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                                                                                                    07-APR-2005
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                                     Synthetic.
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                                                                                                                                                                                                                                          Golz S,
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a subject. (MI) involves administering to the subject, a short interfering nucleic acid (siNA) molecule (I) that directs cleavage of a Fas RNA through RNA interference (RNAI) under conditions suitable to modulate the expression of Fas in the subject. Also described: (I) an expression vector comprising (I); (2) a kit comprising (I); (3) a human cell comprising (I); (4) a pharmaceutical composition comprising (I); and (5) a method of synthesizing (I). The present sequence represents a human Fas ligand (FasL) tumor necrosis factor receptor superfamily member 6 (TNFSFE) target and upper (sense) aiRNA oligonucleotide, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating spinal cord injury in subject, involves administering to subject, short interfering nucleic acid directing cleavage of Fas RNA through RNA interference under conditions suitable to modulate expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method (M1) for treating spinal cord injury in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human FasL TNFSF6 gene lower siRNA sequence SEQ ID NO:508.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19 BP; 17 A; 1 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 33; SEQ ID NO 404; 98pp; English.
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                                                                                                    2002US-0409293P.
2003US-0440129P.
2003WO-US005028.
2003WO-US005346.
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                                                                                       2002US-0408378P
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                                                                                                                                                                                                                                                                                                                                                                            Haeberli P, Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-494870/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fas in subject
                                                                                                  09-SEP-2002;
15-JAN-2003;
20-FEB-2003;
20-FEB-2003;
                                                      06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                            23-MAY-2003;
23-OCT-2003;
                                                                                                                                                               30-APR-2003;
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                                          20-MAY-2002;
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Synthetic.

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STAT-3 siRNA antisense strand, SEQ ID 469.
         17-NOV-2005 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating spinal cord injury in subject, involves administering to subject, short interfering nucleic acid directing cleavage of Fas RNA through RNA interference under conditions suitable to modulate expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.6%; Score 16.4; DB 1; Length 19;
14.4%; Pred. No. 1.1e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 1 A; 0 C; 1 G; 0 T; 17 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 33; SEQ ID NO 508; 98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SIRN-) SIRNA THERAPEUTICS INC.
                                                                                                                                                                                                          2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                               2002US-0362016P.
2002US-0363124P.
2002WO-US015876.
2002US-0386782P.
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2003US-00693059.
2003US-00720448.
2003US-00727780.
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2003WO-US005346.
2003US-00427160.
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2004US-0543480P
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                                                                                      18-JUN-2004; 2004US-00871222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 94.4
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2005-494870/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fas in subject
                            US2005119212-A1
                                                                                                                                                                            20-MAY-2002;
06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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30-APR-2003;
23-MAY-2003;
23-OCT-2003;
24-NOV-2003;
03-DEC-2003;
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20-FEB-2003;
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                                                                                                                                                  06-MAR-2002
                                                        02-JUN-2005
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The invention relates to a novel chemically synthesized double stranded short interfering nucleic acid molecule that directs cleavage of a signal transducer and activator of transcription 3 (STMT3) RNA by RNA interference. The invention further includes a composition comprising the short interfering nucleic acid has cytostatic, antipporiatic, dermatological, antiinflammatory, and gastrointestinal-Gen activities. The short interfering nucleic acid or its composition is useful for treating, preventing, inhibiting, or reducing cancer, proliferative, and/or inflammatory diseases, disorders, or conditions in a subject or organism, such as psoriasis, eczema, dermatitis, Crohn's disease, and inflammatory bowel disease, and for any other disease, trait, or condition that is related to or will respond to the levels of STAT3 in a cell or tissue,
                                                                                                                                                                  gene silencing; cytostatic; antipsoriatic; dermatological; antinflammatory; gastrointestinal-Gen.; cancer; inflammation; psoriasis; eczema; dermatitis; Crohns disease; inflammatory bowel disease; siRNA; short interfering RNA; ss.
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                                                                                                                                               Signal-transducer and activator of transcription-3; RNA interference;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 469; 266pp; English.
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30-ARR-2003; 2003US-00427160.
23-MAY-2003; 2003US-00444853.
23-GC-2003; 2003US-00693059.
24-NOV-2003; 2003US-00720448.
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10-FEB-2004; 2004US-0543480P.
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24-MAY-2004; 2004WO-US016390.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001US-0292217P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-JUL-2001; 2001US-0306883P.
13-AUG-2001; 2001US-0311865P.
20-PEB-2002; 2002US-0358580P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-FEB-2004; 2004US-00780447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2004; 2004US-00826966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-DEC-2004; 2004US-00014373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Robin H, Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2005-604649/62.
                                                                                                                                                                                                                                                                                                                                                                                                   US2005196781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                38-SEP-2005
                                                                                                                                                                                                                                                                                                                                       Synthetic.
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Gaps ö

2705 TACTAAAAAAAAAAA 2722

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94.48;

TACAAAAAAAAAAAA 1

18

)871/c AEC90871 standard; RNA; 19 BP.

RESULT 1407 AEC90871 AEC90871;

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0
                                                                                                                                                                                                                                                                                 Signal-transducer and activator of transcription-3; RNA interference; gene silencing; cytostatic; antipsoriatic; dermatchological; antiinflammatory; gastrointestinal-den.; cancer; inflammation; psoriasis; eczema; dermatitis; Crohns disease; inflammatory bowel disease; siRNA;
alone or in combination with other treatments or therapies. This oligo sequence represents a STAT-3 siRNA strand of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel chemically synthesized double stranded short interfering nucleic
                                                                                  Gaps
                                                                                  ö
                                                          0.6%; Score 16.4; DB 1; Length 19; 94.4%; Pred. No. 1.1e+03;
                                                                                  1; Indels
                                   Sequence 19 BP; 2 A; 0 C; 2 G; 0 T; 15 U; 0 Other;
                                                                                                                                                                                                                                                          STAT-3 siRNA target/sense strand, SEQ ID 192.
                                                                                 0; Mismatches
                                                                                                       2707 CTAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                       BP.
                                                                                                                             CTCAAAAAAAAAAAA 1
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2003US-00720448.
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2002WO-US015876.
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2002US-00201394.
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2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003WO-US005028.
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2002US-0363124P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003US-00427160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-MAY-2004; 2004WO-US016390.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001US-0311865P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004US-0543480P
                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-2004; 2004US-00014373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2004US-00757803
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                                                                                                                                                                                      AEC90594 standard; RNA; 19
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                               short interfering RNA; ss.
                                                                    Local Similarity 94.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2005-604649/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-MAY-2002;
17-MAY-2002;
06-JUN-2002;
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23-OCT-2003;
24-NOV-2003;
03-DEC-2003;
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06-MAR-2002;
11-MAR-2002;
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10-FEB-2004;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                             18
                                                                                                                                                                                                            AEC90594;
                                                           Query Match
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                                                                                                                                                                RESULT 1408
                                                                             Matches
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                                                                                                                           short interfering nucleic acid molecule that directs cleavage of a signal transducer and activator of transcription 3 (97AT3) RNA by RNA interference. The invention further includes a composition comprising the short interfering nucleic acid in a carrier or diluent. The short interfering nucleic acid has cytostatic, antipsoriatic, dermacological, antiinflammatory, and gastrointestinal-Gen activities. The short interfering nucleic acid or its composition is useful for treating, preventing, inhibiting, or reducing cancer, proliferative, and/or inflammatory diseases, disorders, or conditions in a subject or organism, such as psoriasis, eczema, dermatitis, Crohn's disease, and inflammatory bowel disease, and for any other disease, trait, or condition that is related to or will respond to the levels of STAT3 in a cell or tissue, alone or in combination with other treatments or therapies. This oligo sequence represents a STAT-3 siRNA strand of the invention.
acid molecule that directs cleavage of STAT3 RNA through RNA interference, useful for treating cancer and inflammatory diseases e.g.
                                                                                                               The invention relates to a novel chemically synthesized double stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA interference, gene silencing; short interfering RNA; siRNA;
hair disease; depilatory; alopecia; atrichia; endocrine-gen.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.6%; Score 16.4; DB 1; Length 19; 88.9%; Pred. No. 1.1e+03; ve 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human vitamin D receptor siRNA antisense strand, SEQ:451.
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 15 A; 2 C; 0 G; 0 T; 2 U; 0 Other;
                                                                         Example 3; SEQ ID NO 192; 266pp; English.
                                    psoriasis in subject or organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2707 CTAAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JUL-2001; 2001US-0306883P.

13-AUG-2001; 2001US-0306883P.

10-FEB-2002; 2002US-0358580P.

06-MAR-2002; 2002US-0352016P.

11-MAR-2002; 2002US-035124P.

11-MAY-2002; 2002WS-0363124P.

06-JUN-2002; 2002WS-038782P.

29-AUG-2002; 2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 CUCAAAAAAAAAAAAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
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09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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23-MAY-2003; 2003US-00444853
23-OCT-2003; 2003US-00693059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JUL-2004; 2004US-00898311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.98;
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AEE65553 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vitamin D receptor; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2005277608-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-FEB-2006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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The invention relates to chemically synthesized short interfering nucleic acids (siNAs) which downregulate expression of the vitamin D receptor (VDR) gene by RNA interference. The siNAs may or may not comprise ribonucleotides, can contain deoxyribonucleotides, can be chemically conditied and may be double or single stranded. They further comprise coligonucleotide and an antisense regions, or alternatively are assembled from a sense coligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The invention also relates to pharmaceutical compositions comprising an siNA targeted to a human covitamin D receptor manA (see RefSeq accession number NM 000376 (AEB65732)), especially the siRNAs shown in AEB65103-AEB65727. The invention further discloses expression vectors and host cells comprising an siNA of the invention. The siNAs are used to modulate expression of the vitamin D receptor gene in cells, tissue explants or organisms for the treatment of a variety of conditions. siNAs that downregulate vitamin C receptor expression may be used to prevent or reduce hair growth, and D receptor expression may be used to prevent or reduce hair growth, and con be used to targenia e used to prevent or reduce hair growth, and contain D receptor expression can be used in the treatment or prevention of alphoeia (e.g., androgenic alopecia), arricina or other disorder associated with a deficiency of vitamin D receptor expression and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide columns) the antisense strand of a single human vitamin D receptor transcript variant l-targeted double-stranded columns.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                         Novel chemically synthesized double-stranded short interfering nucleic
                                                                                                                                                                                                                                                                                                                                                acid molecule directing cleavage of vitamin D receptor RNA through RNA interference, useful for treating such as alopecia and atrichia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 1 A; 0 C; 1 G; 0 T; 17 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                Claim 33; SEQ ID NO 451; 238pp; English.
                                                                                                                                                                                            (SIRN-) SIRNA THERAPEUTICS INC
                                                        2004US-0543480P.
2004US-00780447.
2004US-00826966.
2004WO-US013456.
                                                                                                                                                 24-MAY-2004; 2004WO-US016390
                  2003US-00727780
                                        2004US-00757803
                                                                                                                                                                                                                                          Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                   WPI; 2006-037965/04.
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nes 17; Conserv
                                                                                                                                                                                                                                       α,
                  03-DEC-2003;
14-JAN-2004;
10-FEB-2004;
                                                                                                      16-APR-2004;
                                                                                 13-FEB-2004;
                                                                                                                                                                                                                                       Guerciolini
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Human vitamin D receptor target sequence/siRNA sense strand, SEQ:195.
2705 TACTAAAAAAAAAAA 2722
                                                                                                                                                                     BP
                                            18 TACAAAAAAAAAAA
                                                                                                                                                                AEE65297 standard; RNA; 19
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RNA interference; gene silencing; short interfering RNA; siRNA;

09-FEB-2006 AEE65297;

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RESULT 1410

ઠ 유 AEE65297

The invention relates to chemically synthesized short interfering nucleic acids (sinAs) which downregulate expression of the vitamin D receptor (VDR) gene by RNA interference. The sinAs may or may not comprise ribonucleotides, can contain deoxyribonucleotides, can be chemically sense and antisense regions, or alternatively are assembled from a sense coligonucleotide. Specifically, the sinAs include short interfering RNA (siRNA). The invention also relates to pharmaceutical compositions comprising an sinA targeted to a human comprasing an sinAs include short interfering RNA (siRNA). The invention also relates to pharmaceutical compositions comprising an sinA targeted to a human comprasing the specially the siRNAs shown in AEBES103-AEE6572. The caspecially the siRNAs shown in AEBES103-AEE6572. The caspecially the siRNAs shown in AEBES103-AEE6572. The caspecially the siRNAs shown in AEBES103-AEE6572. The caspecial of an siRNAs shown in AEBES103-AEE6572. The caspecial of an include expression of the treatment of a variety of conditions. SiNAs that downregulate vitamin D receptor gene in cells, tissue explantes or organisms of the treatment of a variety of conditions. SiNAs that downregulate vitamin D receptor anaphase in hair folicites for hair removal (depilation). SiNAs that act to upregulate vitamin D receptor expression can be used to target anaphase in hair folicites for hair removal (depilation). SiNAs that act to upregulate vitamin D receptor expression con the treatment or prevention con vitamin D receptor expression of an inhibitor of such a slopecia (e.g., androgenic alopecia), articitia or other disorder associated with a deficiency of vitamin D receptor expression. The siNAs Novel chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of vitamin D receptor RNA through RNA interference, useful for treating such as alopecia and atrichia. hair disease; depilatory; alopecia; atrichia; endocrine-gen.; Claim 33; SEQ ID NO 195; 238pp; English. (SIRN-) SIRNA THERAPEUTICS INC 2001US-0311865P. 2002US-0358580P. 2002US-0362016P. 2002US-0363124P. 2002WO-US015876. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 2003US-0440129P. 2003WO-US005028. 2003WO-US005346. 2003US-00417012. 2003US-00427160. 2003US-00444853. 23-JUL-2004; 2004US-00898311 2002US-0386782P. 2003US-00693059. 2003US-00720448 2003US-00727780 2004US-00757803 2004US-0543480P 30-APR-2004; 2004WO-US013456. 24-MAY-2004; 2004WO-US016390. 2004US-00780447 Mcswiggen J; vitamin D receptor; ss WPI; 2006-037965/04. US2005277608-A1 Guerciolini R, 29-AUG-2002; 2 05-SEP-2002; 2 09-SEP-2002; 2 20-FEB-2003; 16-APR-2003; 11-MAR-2002; 17-MAY-2002; 06-JUN-2002; 20-FEB-2003; 30-APR-2003; 23-OCT-2003; 24-NOV-2003; 03-DEC-2003; Homo sapiens. 06-MAR-2002; 15-JAN-2003; 23-MAY-2003; 14-JAN-2004; LO-FEB-2004; L6-APR-2004; 

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                                                                                                                                                                                                                                                                                                                                                                    RNA interference; gene silencing; short interfering RNA; siRNA; cancer; neoplasm; hyperproliferation; cytostatic; respiratory disease; respiratory-gen.; cardiovascular disease; cardiovascular-gen.; cottlar disease; ophthalmological; diabetic retinopathy; antidiabetic; stromal cell-derived factor-1; SDF-1; chemokine CXC motif ligand 12;
may also be used in drug screening, diagnostics, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the sense strand of a human vitamin D receptor transcript variant 1-targeted double-stranded siRNA, which is identical to the human vitamin D receptor transcript variant 1 target sequence.
                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                              Human SDF-1 (CXCL12b) target sequence/siRNA sense strand, SEQ:395
                                                                                                                                                        ;
                                                                                                                            Score 16.4; DB 1; Length 19;
Pred. No. 1.1e+03;
1; Mismatches 1; Indels
                                                                                                    Sequence 19 BP; 17 A; 1 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                             2705 TACTAAAAAAAAAAA 2722
                                                                                                                                                                                                       UACAAAAAAAAAAA 19
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2003MO-US005346.
2003US-00444853.
2003US-0059059.
2003US-00727408.
2003US-007277803.
2004US-00757803.
2004US-00757803.
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2002WG-US0138124P.
2002WG-US018F076.
2002US-0406784P.
2002US-040878P.
2002US-0408378P.
2003US-0408293P.
                                                                                                                                                                                                                                                                   AEF36928 standard; RNA; 19 BP
                                                                                                                             0.6%;
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                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                         Query Match
Best Local Similarity 88.9°
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2006019917-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAR-2002;
11-MAR-2002;
17-MAY-2002;
06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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20-FEB-2003;
20-FEB-2003;
30-APR-2003;
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10-FEB-2004;
13-FEB-2004;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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23-OCT-2003;
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03-DEC-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    CXCL12; ss.
                                                                                                                                                                                                                                                                                             AEF36928;
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The invention relates to chemically synthesized short interfering nucleic acids (siNAs) which downregulate stromal cell-derived factor-1 (SDF-1, chemokine CXC motif ligand 12, CXCL12) gene expression by RNA interference. The siNAs are characterized in that one or more nucleotides is chemically modified to reduce the immunostimulatory properties of each sinA to a level below that of a corresponding unmodified siNA.

CC siNA to a level below that of a corresponding unmodified siNA.

CC sinAt to a level below that of a corresponding unmodified siNA.

CC ditther comprise sense and antisense regions, or alternatively are contain deoxyribonucleotides, and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are contain a sense oligonucleotide and an antisense oligonucleotide.

CC further comprise sense and antisense regions, or alternatively are specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA), the siNAs include short interfering RNA (siRNA). The siNAs of the invention are used to condulate expression of the SiP-1 gene in cells, tissue explants or corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants correction of cancer and other proliferative conditions, respiratory diseases, cardiovascular diseases and coular diseases, and are especially cuseful for the treatment of fabetic retinopathy (proliferative cretinoplasty). The siNAs may also be used in drug screening, therapeutic target identification and validation, genetic engineering, parameted conditions, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the construction and validation in degeneral conditions, the sinAs may also be used in drug screening, settled to the sinAs may also be used in drug screening.

CC stanscript variant 1 (CXCL12b), which is identical to the human SDF-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                   molecule directing cleavage of stromal cell derived factor-1 RNA, useful for treating cancer and ocular, respiratory and cardiovascular diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA interference, gene silencing, short interfering RNA, siRNA, cancer, neoplasm, hyperproliferation, cytostatic, respiratory disease, respiratory-gen., cardiovascular disease, cordiar disease, ophthalmological, diabetic retinopathy, antidiabetic, stromal cell-derived factor-1, SDF-1, chemokine CXC motif ligand 12;
                                                                                                                                       Chemically synthesized double-stranded short interfering nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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4%; Pred. No. 1.1e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human SDF-1 (CXCL12b) siRNA antisense strand, SEQ:574.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 16 A; 3 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                      Example 3; SEQ ID NO 395; 384pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2706 ACTAAAAAAAAAAAA 2723
(SIRN-) SIRNA THERAPEUTICS INC.
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nes 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AEF37107 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variant 1 target sequence.
                                                                                            WPI; 2006-134231/14.
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AEF37107/c
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transcript variant 1 (CXCL12b).
                                                                                                 WPI; 2006-134231/14.
                                                                                            Guerciolini R,
                             29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                           20-FEB-2003;
                     11-MAR-2002;
17-MAY-2002;
                                     15-JAN-2003;
20-FEB-2003;
                                                23-MAY-2003;
                                                           14-JAN-2004;
                  06-MAR-2002;
                          06-JUN-2002;
                                                        03-DEC-2003;
                                                                            20-AUG-2004
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Length 19; Score 16.4; DB 1; Length 1 Pred. No. 1.1e+03; 0; Mismatches 1; Indels 2706 ACTAAAAAAAAAAAA 2723 Claim 4; Page 29; 64pp; English. BP. 97WO-CA000454. 0.6%; 96US-0021152P ACCAAAAAAAAAAAAAA AAV12302 standard; DNA; 20 17-JUN-1998 (first entry) Query Match
Best Local Similarity 94.4
Matches 17; Conservative Wright JA, Young AH; WPI; 1998-086958/08. (WRIG/) WRIGHT J A. (YOUN/) YOUNG A H. 01-JUL-1996; 30-JUN-1997; Homo sapiens WO9800532-A2 08-JAN-1998. AAV12302; 18 RESULT 1413 AAV12302 ò 셤 The invention relates to chemically synthesized short interfering nucleic acids (siNAs) which downregulate stromal cell-derived factor-1 (SDF-1, CCCL12) gene expression by RNA chemically modified to reduce the immunostimulatory broperties of each sink to a level below that of a corresponding unmodified siNAs.

Cofficially, the siNAs may or may not comprise ribonucleotides, can contain deoxyribonucleotides, and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The stranded that micro-RNA (miRNA) and short hairpin RNA (siRNA). The contained sor relates to pharmaceutical compositions comprising an siNA trargeted to an SDF-1 mRNA. The siNAs of the invention are used to condulate expression of the SDF-1 gene in cells, tissue explants or contained sort the transplants of corganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of avriety of conditions. They may be used in the treatment of ancer and other proliferative conditions, respiratory diseases, and are especially cusful for the treatment of diabetic retinopathy (proliferative cusful for the treatment of diabetic retinopathy (proliferative cusful for the treatment of diabetic retinopathy (proliferative cusing, pharmacogenomics, studying gene function, and gene mapping (e.g., of antisense strand of a double-stranded siRNA targeted to human SDF-1 contined and siRNA targeted to human SDF-1 Chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of stromal cell derived factor-1 RNA, useful for treating cancer and ocular, respiratory and cardiovascular diseases. Example 3; SEQ ID NO 574; 384pp; English. (SIRN-) SIRNA THERAPEUTICS INC 2003WO-US005346. 2003US-00444853. 2003US-00643059. 2003US-00720448. 2003US-00727780. 2004US-00757803. 2004US-00757803. 2002WO-US015876. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 2003US-0401298. 2003WO-US005028. 2004WO-US016390. 2004US-00923536. 2002US-0362016P 2002US-0363124P 04-APR-2005; 2005US-00098303 27-MAY-2005; 2005US-00140328 Mcswiggen J;

Sequence 19 BP; 0 A; 0 C; 3 G; 0 T; 16 U; 0 Other;

The present sequence represents a 3'-untranslated region (3'UTR) fragment of ribonucleotide reductase R1. The present invention describes: (1) cof ribonucleotides (ON) comprising at least 7 consecutive nucleotides (IN) comprising at least 7 consecutive nucleotides (IN) conjudence on UTR of a housekeeping gene; (2) antisense ON CC (AON) complementary to ON; (3) ribozymes (Rb) complementary or homologues (AON) complementary to ON; (3) ribozymes (Rb) complementary or homologues (CC (5) an antibody (Ab) that binds to ON, OAN and Rb; (6) ant probe in P CC (5) an antibody (Ab) that binds to ON, OAN and Rb; (6) ant probe in P CC (5) an antibody (Ab) that binds to ON, AON, AON, Rb and Ab are used to modulate (especially inhibit) growth of tumour cells (especially neoplastic cells) and to reduce their capacity for metastasis. The above may also be used to treat benign proliferative disorders e.g. human immunodeficiency virus, cangiogenesis and viral infections, e.g. human immunodeficiency virus, hepatitis or herpes. ON may further be used: (1) to identify modulators of tumour growth/metastasis; (ii) to identify compounds (especially compounds (especially for monitoring properses of cancer therapy. SON inhibit tumour gentification and quantification compounds of cumorification cells, particularly where these are ö ö New oligo-nucleotide(s) complementary to untranslated regions of housekeeping genes - are useful in, e.g. identifying modulators of tumour growth/metastasis and inhibiting growth of neoplastic cells. Ribonucleotide reductase R1; 3'-untranslated region; 3'UTR; tumour; housekeeping gene; identification; modulator; metastasis; neoplastic; papilloma; atherosclerosis; angiogenesis; viral infection; ss. Gaps Gaps . 0 ; 0.6%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 1.18+03; tive 0; Mismatches 1; Indels Ribonucleotide reductase R1 3'UTR fragment SEQ ID NO:46. Sequence 20 BP; 17 A; 1 C; 2 G; 0 T; 0 U; 0 Other; Best Local Similarity 94.4 Matches 17; Conservative Query Match 

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WO200105435-A2.
                    25-JAN-2001.
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                                                                                                                                      Human; fucosyltransferase; FUT3; FUT6; cancer; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                              The present invention provides antisense oligonucleotides to the human fucosyltransferase coding sequences, particularly FUT3 and FUT6. These antisense sequences can be used in the treatment of cancer, especially colon, pancreatic, ovarian, gastric, breast, lung, hepatocellular, prostate, bladder, renal cell and uterine cancers. In addition, they can also be used in the treatment of animals such as dogs, cats and horses
                                                                                                                                                                                                                                                                                                                  Novel antisense human fucosyltransferase sequences useful for treating cancer including breast, lung, gastric, renal and uterine cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Insulin-like growth factor binding protein-5; IGFBP-5; human;
antisense oligonucleotide; hormone-regulated cancer; prostatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.6%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 1.1e+03; ve 0; Mismatches 1; Indels
                                                                                                                    Human FUT6 antisense oligonucleotide SEQ ID NO: 19.
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2565 TCTCCTGAGCTAGGAAGA 2582
                                                                                                                                                                                                                                                                                                                                              Claim 7; Page 33; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense IGFBP-5 inhibitor #13.
                  AAAAAAGAAAAAAAA 18
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                                                               BP
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                                                                                                                                                                                                                                           99US-0131068P
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                                                                                                                                                                                                                                                            (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.48;
                                                             AAC68768 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA91207 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                breast cancer; therapy; ss.
                                                                                                 20-FEB-2001 (first entry)
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Matches 17; Conservative
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                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                   WO200064262-A1.
                                                                                                                                                                  Homo sapiens.
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2709
                                                                                 AAC68768;
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                                            RESULT 1414
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This sequence represents an antisense oligonucleotide targeted against human insulin-like growth factor binding protein-5 (IGFBP-5). The invention relates to a composition for treatment of hormone-regulated cancer, comprising an antisense oligonucleotide (such as this sequence) which inhibits expression of IGFBP-5 by hormone-regulated tumour cells. The compositions is useful for delaying progression of hormone-regulated tumour cells such as prostatic cancer cells or breast cancer cells, to an antrogen-independent state, by treating hormone sensitive tumour cells androgen-independent state, by treating hormone sensitive tumour cells cumour cells. The composition can also be used for treating a hormone-responsive cumour cells in individual and administering the composition to the individual after initiation of hormone-withdrawal to induce apoptotic cell death of hormone-responsive tumour cells, and therefore delaying the progression of hormone-responsive cancer cells to a hormone-independent composition to individual. It can also be used for inhibiting or delaying metastatic boney progression of an IGF-1 sensitive tumour in a mammal, by administering the composition to inhibit the expression of IGFBP-5 by the composition to inhibit the expression of delaying content of the composition to inhibit the expression of delaying content of the content of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Composition for treating hormone-regulated cancer, e.g. breast and prostatic tumors, comprising an antisense oligonucleotide that inhibits expression of insulin like growth factor binding protein-5 by hormone-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 34; 45pp; English.
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                                                                                                                                                                                                                              (UYBR-) UNIV BRITISH COLUMBIA.
(MIYA/) MIYAKE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGAAAAAAAAAAAAA
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                                                                                                                     99US-0144495P
19-JUL-2000; 2000WO-CA000853
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                                                                                                                     19-JUL-1999;
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                                                                                                                                                    The invention relates to a novel polynucleotide isolated and purified from a human gene having any one of 935 fully defined sequences as given in specification, or a sequence having a base substitution. The invention further relates to: an oligonucleotide containing single nucleotide polymorphisms; a PCR primer set chosen from the combination of two DNA fragments from any one of 1220 fully defined sequences as given in specification; a labelling probe containing the SNP containing oligo, and a microarray equipped with the SNP containing the single nucleotide gene of the invention is useful for detecting the single nucleotide
                                                                                            Novel polynucleotide useful for detecting single nucleotide polymorphisms
                                                                                                                                                                                                                                                              polymorphisms in human gene. The isolated human gene is also useful for diagnosis of disease and determination of side effect to a medical agent. The isolated human gene is also effective in detecting single nucleotide polymorphisms in a human gene. This polymucleotide sequence represents one of the PCR primers used in the single nucleotide polymorphism detection method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human, antisense; lung dysfunction; nasal airway dysfunction; antisifammatory steroid; ubiguinone; antisifammatory; antisilergic; antisathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                                                                                                                                                                                                                                                                                                          0.6%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 6 A; 8 C; 3 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inflammation; respiratory disease; ds.
                                                                                                                                 Claim 2; SEQ ID NO 1519; 704pp; Japanese.
                                               (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sandrasagra A, Ke
L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                          2210 GGCTCCTTGGTGGATGAG 2227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ85532 standard; DNA; 20 BP.
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12-FEB-2002; 2002JP-00034717
                       12-FEB-2002; 2002JP-00034717
                                                                                                                                                                                                                                                                                                                                                                                                                                             GGCTTCTTGGTGGATGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                  17; Conservative
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Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-229219/22
                                                                       WPI; 2003-820454/77
                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200285308-A2
                                                                                                           in human gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ85532;
                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1417
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' end advinctions of genes encoding prepride associated with lung and/or junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention comparation and a second active agent comprising an antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition may have a composition at sepiratory, lung or mallignamt disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an entiinflammatory steroid in a subject, for reducing or depleting levels of contining sensitivity to adenosine, reducing or depleting levels of continion, continuation, lung allergies, or a respiratory disease or condition, lung allergies, or a respiratory disease or condition.

Conting inflammation, lung allergies, or a respiratory disease or condition, and the printed contined in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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Pharmaceutical composition for treating ailments associated with impaired
                                      respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 1.1e+03;
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                                                                                                                                                                                   Claim 15; SEQ ID NO 774; 872pp; English.
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Tang L, Shahabuddin S;
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Miller S,
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25-MAR-2004
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating construction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, curfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery of device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production or reating a respiratory, lung or mailgnant disease. The administered cor availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and asease or condition such as pulmonary vasoconstriction, inflammation, allergies and/or surfactant hypoproduction are associated with a mannarion and asease.
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                                                                                                                                                                                                     ဌ
                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted trucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 774; 763pp; English
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WPI; 2003-093058/08.
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                              0.6%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 1.1e+03;
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Sequence 20 BP; 17 A; 2 C; 1 G; 0 T; 0 U; 0 Other;
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es 17; Conservative
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2724 2707 CTAAAAAAAAAAAAA з ссададададададада

380/c ADH66380 standard; DNA; 20 BP. (first entry) ADH66380; RESULT 1419

Human glucocorticoid receptor-specific antisense oligonucleotide #3214.

antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.

Homo sapiens

4O2003099215-A2

04-DEC-2003

20-MAY-2003; 2003WO-US016084.

20-MAY-2002; 2002US-0381857P.

(PHAA ) PHARMACIA CORP

Crosby SD, Nalseth AE;

WPI; 2004-035034/03.

New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.

Claim 4; SEQ ID NO 3214; 985pp; English.

to nucleic acids encoding a mammalian glucocorticoid receptor. The antiense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone. The invention comprises an antisense oligonucleotides that are

Sequence 20 BP; 2 A; 1 C; 1 G; 16 T; 0 U; 0 Other;

Gaps ô 0.6%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 1.1e+03; ve 0; Mismatches 1; Indels 94.48; Conservative Local Similarity les 17; Conserv Query Match Matches

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2708 TAAAAAAAAAAAAAA 2725 TCAAAAAAAAAAAAAAA

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BP. ADJ61530 standard; DNA; 20 (first entry) 06-MAY-2004 ADJ61530; RESULT 1420 ADJ61530 THE STANDARD STANDARD

Oligonucleotide associated to IL5R-X61176 #222.

interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;

Homo sapiens

WO2004011613-A2.

05-FEB-2004.

25-JUL-2003; 2003WO-US023509.

29-JUL-2002; 2002US-0399076P.

(EPIG-) EPIGENESIS PHARM INC

Nyce JW, Tang L, Sandrasagra A, Aguilar D, Miller S; Shahabuddin S, Lu H, Cong H;

WPI; 2004-203534/19.

e.g., Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory

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The present invention relates to an oligonuclectide anti-sense to e.g., initiation codon, coding region with 2-10 nuclectides of 5'-end and 3'-end of nucleic acid trarget comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the cidentellar optionally surfactant operatively linked to the oligonuclectide and optionally surfactant operatively linked to the coligonuclectide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of subject an effective amount of an inhibitor. The oligonuclectide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ses), asthma, impeded chosen from airway inflammation, costic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric phosphorothioate oligonucleotide to target Navl.3 #1059.
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Pred. No. 1.1e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 6 A; 3 C; 5 G; 6 T; 0 U; 0 Other;
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                                     Claim 2; SEQ ID NO 2386; 85pp; English.
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Best Local Similarity 94.4%;
Matches 17; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTTGCTGAGGATAAAAG
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disease e.g., asthma
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                                                                                                                                                                                                                                                                                                                                                                                        invention.
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associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphororhioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Ectaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; are airway inflammation; allergy; impeded respirators, cystic fibrosis; chronic obstructive pulmonary disease; cOPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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                                                                                                                                                                                Sequence 20 BP; 1 A; 2 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                Score 16.4; DB 1;
Pred. No. 1.1e+03;
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                                                                                                                                                                                                                               94.4%; Pred. ww. rive 0; Mismatches
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Cong H;
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                                                                                                                                                                                                                     0.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human oligonucleotide #2286.
                                                                                                                                                                                                                                                                                                                                                                                               ADO46920 standard; DNA; 20
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                                                                                                                                                                                                                               Best Local Similarity 94.4
Matches 17; Conservative
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SANDRASAGRA A.
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SHAHABUDDIN
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AGUILAR D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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(SAND/)
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(SHAH/)
(LUHH/)
(CONG/)
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invention.
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The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target chosen from a gene encoding interleukin (IL) -4 receptor, interleukin (IL) -5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The prevention and/or treatment of a respiratory or lung disease. The gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonuclectides are useful for reating a respiratory or lung disease. The respiratory or lung disease is associated with hyper-responsiveness to receptor of a medical control of a denosine and/or levels of denosine A receptor(8), and/or asthma and/or lung allergies associated with inflammation or an inflammatory disease. The respiratory or lung disease is chosen from airway inflammation, allergy, asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD), allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of the

Sequence 20 BP; 6 A; 3 C; 5 G; 6 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 16.4; DB 1; Length 20; larity 94.4%; Pred. No. 1.1e+03; Conservative 0; Mismatches 1; Indels CTTTGCTGATGATAAAG 1986 CTTTGCTGAGGATAAAG 18 Local Similarity les 17; Conserv 1969 -Query Match Matches ò 셤

RESULT 1423

ADP69379 standard; DNA; 20 (first entry) 09-SEP-2004 ADP69379; 

ВP

Human mitoNEET-specific antisense oligonucleotide #273.

human; antisense oligonucleotide; mitochondrial membrane; insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.

Homo sapiens.

WO2004053060-A2.

24-JUN-2004.

25-NOV-2003; 2003WO-US037621.

06-DEC-2002; 2002US-0431529P.

(PHAA ) PHARMACIA CORP.

Colca JR;

WPI; 2004-468836/44.

New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.

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The invention comprises antisense oligonucleotides that are targeted to membranes, which bind insulin sensitising, antidiabetic thiazolodinediones (referred to as: mitoNEET). The antisense oligonucleotides of the invention are useful for modulating mitoNEET expression and for treating diseases or conditions associated with mitoNEET, such as: diabetes, immunological disorders, cardiovascular disorders including hypertension, neurological disorders, and sischaemia/reperfusion injuries. The present DNA sequence represents a mitoNEET-specific antisense oligonucleotide of the invention. NOTE: The phosphorothioate backbone.
Claim 4; SEQ ID NO 273; 226pp; English.
  8 X C C C C C C C C C C X 8
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Sequence 20 BP; 3 A; 0 C; 1 G; 16 T; 0 U; 0 Other;

Gaps ö Score 16.4; DB 1; Length 20; Pred. No. 1.1e+03; 1; Indels 0; Mismatches 0.6%; 94.48; 17; Conservative Similarity Query Match Local Matches

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2708 TAAAAAAAAAAAAAA 2725 TAAACAAAAAAAAA 1 18 ð 셤

AAX18389 standard; DNA; 18 RESULT 1424 AAX18389/c

BP. (first entry) 11-MAY-1999 AAX18389; 

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RT-PCR primer of the invention SEQ ID 30.

RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.

Synthetic.

JP11032765-A. 09-FEB-1999.

97JP-00208312. 18-JUL-1997; 97JP-00208312. 18-JUL-1997;

(TAKI ) TAKARA SHUZO CO LTD.

WPI; 1999-183822/16.

Peptides having at least two new nucleotides - useful as primers in RT-

Example 1; Page 12; 19pp; Japanese.

to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta-N3'; or (X)m5'-(gamma)k-delta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine, cytosine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma; in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene This sequence represents a primer of the invention. The invention relates sednences

Sequence 18 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 2 Other;

Query Match

0.6%; Score 16.2; DB 1; Length 18;

Secreted protein; fetal kidney; ds.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta-N3'; or (X)m5'-(agmma) K-delta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thywine; n = netural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine or cytosine; N = adenine, guanine or thymine; k = netural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1;3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides having at least two new nucleotides - useful as primers in RT-
                                                                                                                                                                                                                                                                              RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.
                           Gaps
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                           Indels
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94.1%; Pred. No. 1.1e+03;
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100.0%; Pred. No. 1.1e+03;
:ive 0; Mismatches 0;
                                                                                                                                                                                                                                                RT-PCR primer of the invention SEQ ID
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                                                          2708 TAAAAAAAAAAAAA
                                                                                     17 BAAAAAAAAAAAAAAA
                                                                                                                                                            AAX18368 standard; DNA; 16
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Les 16, Conservative
                             16; Conservative
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             Best Local Similarity
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                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                  RESULT 1425
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                              Matches
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The sequence is that of the 3' end of a sequence encoding a secreted protein from a human fetal kidney clone AK296. Such a sequence is predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals, although no supporting data is given. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immacopolesis regulating (e.g. as proliferation/differentiation activity, haemacopolesis regulating activity, tissue growth activity, activity, haemacopolesis regulating activity, chemotactic/chemokinetic activity, haemacopolesis and thrombolytic activity, inceptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity. It is also stated to be useful for gene therapy
                                                                                                                                                                                                                                                                  New polynucleotides encoding secreted human proteins - are derived from foetal kidney or adult retina cDNA libraries, used as, e.g. potential
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                                                                                                                                                                                               Merberg D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 16; DB 1; Length 16; 00.0%; Pred. No. 1.1e+03; ve 0; Mismatches 0; Indels
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                                                                                                                                                                                               Lavallie ER, Racie LA,
                                                                                                                                                                                                                                                                                                                               Disclosure; Page 54; 76pp; English
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Best Local Similarity 100.0%; Pr
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28-JAN-2000; 2000DE-01003631
                                                                                                                                         97US-00885610
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                                                                                                                                                                  (GEMY ) GENETICS INST INC.
                                                                                                                                                                                               Mccoy JM, I
Agostino MJ;
                                                                                                                                                                                                                                          WPI; 1999-095671/08.
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                                                                                                                                         30-JUN-1997;
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                                                      WO9900405-A1
                           Homo sapiens
                                                                                 07-JAN-1999.
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                                                                                                                                                                                                                                                                                                     vaccines.
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This invention describes nucleoside derivatives (I) with photolabile protecting groups. (I) are used to synthesize oligonucleotides using the photolithographic nucleic acid chip method, particularly where these are intended for performing enzymatic reactions initiated from a free 3. hydroxy (especially solid-phase polymerase reactions or ligase reactions, but also reverse transcription, cDNA synthesis etc.), also for hybridization testing, sequencing and in DNA computing. (I) are produced with high selectivity by reaction with a mild acylating agent that has high specificity for the 3. position, without significant side-reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "OTHER = F1(CH2)6-PO-thymine, where F1 is flavine and PO is a phosphate group"
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                                                                                                                                                                                                                 New nucleoside derivatives with photolabile protecting groups, use oligonucleotide synthesis, particularly on solid phases, e.g. for hybridization testing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analytical support; genomic sequencing; mutation detection; pharmaceutical development; ss.
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                             (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM
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100.0%; Pre-
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                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 9; 48pp; German.
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                                                                                             Hoheisel J;
                                                                                                                                                    WPI; 2000-679457/66.
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modified_base
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                                                                                             Beier M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABA04585;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1428
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Matches
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Decout JL,

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The present sequence is that of an oligonucleotide (ODN)-minor groove binder (MGF) complex. MGBs bind in a non-intercalating manner to the minor groove of non-single-stranded DNA, RNA or their hybrids. ODN-MGB-LF conjugates of the invention also comprise a latent fluorophore (LF), which binds similarly to the MGB but in an intercalating manner, or lies in the minor groove, or is oriented in some other way to the DNA molecule by MGB, such that it becomes fluorescent (or its fluorescent properties change detectably). The conjugates are used as hybridisation probes and
                                                                The present invention relates to an analytical support, to which a number of Oligonucleotides are labelled with a fluorescent compound, the fluorescence of which varies when the oligonucleotide hybridises to its complement. The analytical support is useful in hybridisation testing for identification of specific nucleic acids, such as genomic sequencing, detecting mutations or pharmaceutical development. The present oligonucleotide was used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conjugate of oligonucleotide, minor groove binder and latent fluorophore, useful for detecting specific nucleic acids, e.g. for single-nucleotide mismatch discrimination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ODN-MGB-LF; oligonucleotide; minor groove binder; latent fluorophore; hybridisation; detection; fluorescence; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag= a
/note= "thymine modified by a minor groove binder (2-
dimethylaminonaphthalene-6- sulfonamide"
                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                       0.6%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                     Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
 the fluorescence changes follow hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide-minor groove binder complex.
                                                                                                                                                                                                                                                                100.0%; Pred. wc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vermeulen NMJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 101; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                 Example 1; Page 12; 33pp; French.
                                                                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPOC-) EPOCH BIOSCIENCES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00428236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-OCT-2000; 2000WO-US029786.
                                                                                                                                                                                                                                                                                                                                                            16 AAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF30895 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 16; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ODN-MGB-LF;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF30895;
                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1429
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Gaps

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Score 16; DB 1; Length 16; Pred. No. 1.1e+03; 0; Mismatches 0; Indels

BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;

Sequence 16

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The present sequence is that of the oligonucleotide (ODN) component of an ODN-MGB (minor groove binder)—LF (latent fluorophore) conjugate of the invention. MGBB bind in a non-intercalating manner to the minor groove of invention. MGBB bind in a non-intercalating manner to the minor groove of the on-single-stranded DNA, RNA or their hybrids, while a LF binds similarly but in an intercalating manner, or lies in the minor groove, or is confected in some other way to the DNA molecule by MGB, such that it conjugates are used as hybridisation probes and amplification primers or for fluorescent detection of specifically hybridising sequences, for analysis or diagnosis, especially (real-time) PCR, for single-nucleotide mismatch discrimination, target or signal amplification, array-based assays and sequencing, including detection of double-stranded DNA by triplex formation. Many different targets can be detected a single reaction-triggered fluorescence. Upon hybridisation to the complementary target sequence there was an increase in fluorescence complementary target sequence there was an increase in fluorescence complementary target sequence there was an increase in fluorescence between the ODN-MGB-LF conjugate and its target sequence to the fluorescence emitted by unhybridised (i.e. single-stranded) ODN-MGB-LF, of 8.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conjugate of oligonucleotide, minor groove binder and latent fluorophore, useful for detecting specific nucleic acids, e.g. for single-nucleotide mismatch discrimination.
amplification primers for fluorescent detection of specifically hybridising sequences, for analysis or diagnosis, especially (real-time) PCK, for single-nucleotide mismatch discrimination, target or signal amplification, array-based assays and sequencing, including detection of double-stranded DNA by triplex formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ODN-MGB-LF; oligonucleotide; minor groove binder; latent fluorophore; hybridisation; detection; fluorescence; probe; ss.
                                                                                                                                                                                                  Gaps
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                                                                                                                                                          Length 16;
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                                                                                                                     Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                          DB 1; Le
1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide portion of ODN-MGB-LF conjugate.
                                                                                                                                                          0.6%; Score 16; DB
100.0%; Pred. No. 1.1
:ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 58; 105pp; English.
                                                                                                                                                                                                                                         2709 AAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                             AAF30880 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPOC-) EPOCH BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00428236
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                                                                                                                                                                                                                                                                           16 AAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                        Query Match
Best Local Similarity 100.º
Matches 16, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-328656/34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                   AAF30880;
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                                                                                                                                                                                                                                                                                                                                        RESULT 1430
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moieties optionally extended by a polymerase. The branched chain compounds of the invention are used in nucleic acid synthesis reaction, primer extension reaction, reverse transcription reaction of RNA into DNA, nucleic acid hybridization experiment (for identifying sequence of a nucleic acid), and nucleic acid amplification experiment (for analysing the expression pattern of genes). The compounds are also used in solid-phase enzymatic reactions. The present sequence was used in the course of the invention to produce branched chain compounds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The specification describes branched compounds containing nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gapa
                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Branched compounds useful in e.g. nucleic acid synthesis reaction comprises nucleic acid moieties optionally extended by a polymerase.
                                                                                                                                                                                                                                                             Branched chain compound; nucleic acid synthesis; primer extension; reverse transcription; nucleic acid hybridization; nucleic acid amplification; 88.
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100.0%; Pred. No. 1.1e+03;
ive 0; Mismatches 0; Indels
 Length 16;
                                                                                                                                                                                                                                     Oligonucleotide used to produce branched chain compounds.
                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
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0
0.6%; Score 16; DB 1; Le
100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schmidt W, Hiller R, Huber M, Mueller M;
                                                                                                                                                                                                                                                                                                                                                                                                                      c
"branch present"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            b
"COOH attached"
                                                                                                                                                                                                                                                                                                                                                                               a
"COOH attached"
                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                          2709 AAAAAAAAAAAAAA 2724
                                                                                                                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99EP-00125484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99EP-00125484
                                                                                                                                                        AAH42481 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LION-) LION BIOSCIENCE AG. (VBCG-) VBC GENOMICS GMBH.
                                                                                                                                                                                                             (first entry)
                                                                              16 AAAAAAAAAAAAAA
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Best Local Similarity 100.
Matches 16; Conservative
                                Conservative
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/note=
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/note=
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/note=
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                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
                                                                                                                                                                                                                                                                                                                                                                modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                         misc feature
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                                                                                                                                                                                                             01-OCT-2001
                                16;
                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                   AAH42481;
     Query Match
                                                                                                                              RESULT 1431
AAH42481/c
                                Matches
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2'F-ANA antisense oligo #6, to elicit RNase H degradation of target RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel acyclic linker-containing oligonucleotide useful for preventing decreasing translation, reverse transcription and/or replication of a target RNA in a system, comprises a modified deoxyribonucleotide.
                                                                                 Acyclic linker; gene expression; gene therapy; ribonuclease; RNase H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Fig 7; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                        29-OCT-2002; 2002WO-CA001628.
                                                                                                                                                                                                                                                                                                                                                                                                       29-OCT-2001; 2001US-0330719P.
                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Damha MJ, Viazovkina
                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYMC-) UNIV MCGILL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-421516/39.
                                                                                                                                                                                                                                                                                                                    WO2003037909-A1.
                                                                                                   antisense; ss.
                                                                                                                                                          Key
modified base
                                                                                                                             Unidentified
                                                                                                                                                                                                                             misc_feature
                             07-AUG-2003
                                                                                                                                                                                                                                                                                                                                                08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
 AAD56451;
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This invention relates to oligonuclectide analogues comprising a protein nucleic acid molecule (PNA) monomer. They are used in the detection and separation of nucleic acid molecules and as probes, primers, linkers, adapters and antisense agents on solid supports. Modifications enhance their use as capture and detection probes e.g. by the incorporation of biotin, digoxigenen, radioisotopes, fluorescent labels such as Liuorescein and reporter molecules such as alkaline phosphatase. They are also used for enhancing or inhibiting the activity of an enzyme or cellular activity. The compounds are stable to nucleases and proteases, have high affinity, binding specificity and solubility. The polyamide backbone of FNAs is resistant to both nucleases and proteases. PNAs bind nucleic acid molecules with greater affinity than DNA or RNA concentration. The compounds are relatively simple to synthesize and are used in a wide variety of applications. This sequence represents a DNA coligoner which is used to represent the thermal stability of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     separation and purification dimers and oligomers.
                                                                                                                                                                                                 Nucleotide sequence of oligomer # 1 used to test thermal stability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 16; DB 1; Length 16;
100.0%; Pred. No. 1.1e+03;
lve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Efimov V, Fernandez J, Archdeacon D, Archdeacon J;
Chakhmakhcheau O, Buryakova A, Choob M, Hondorp K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotides analogs useful in detection, of nucleic acid molecules, comprise monomers,
                                                                                                                                                                                                                             Protein nucleic acid molecule; PNA; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 17; Page 118; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; FEE
             2709 AAAAAAAAAAAAA 2724
                                                                                                              ABA97402 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                         14-MAR-2000; 2000US-0189190P.
30-NOV-2000; 2000US-0250334P.
                                                                                                                                                                                                                                                                                                                                              13-MAR-2001; 2001WO-US008111.
                                      16 AAAAAAAAAAAA 1
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                                                                                                                                                                     (first entry)
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                   (ACTI-) ACTIVE MOTIF.
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                                                                                                                                                                                                                                                                                    WO200168673-A1.
                                                                                                                                                                     18-JUN-2002
                                                                                                                                                                                                                                                                                                                  20-SEP-2001
                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                          ABA97402;
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Min

Parniak MA,

Mangos MM,

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/\*tag= b /note= "Bases 8 and 9 are linked by two secouridine linkers which is represented as S in page 49 and X in page 57 and Fig 7 and 8 of the specification"

/note= "2'-deoxy-2'-fluoroarabinothymidine"

mod\_base= OTHER

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Location/Qualifiers

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The invention relates to an acyclic linker-containing oligonucleotide comprising at least one modified deoxyribonucleotide. Oligonucleotides of the invention are useful for preventing or decreasing translation, reverse transcription and/or replication of a target RNA in a system. They are useful for selectively preventing gene expression in a sequencespecific manner, for hybridising to complementary RNA such as cellular mRNA or viral RNA, to hybridise to complementary RNA such as cellular mRNA or viral RNA, to hybridise to and induce cleavage of complementary RNA. They are also useful therapeutically in formulations or medicaments to prevent or treat a disease characterised by the expression of a particular target RNA. The invention is used in gene therapy. The present sequence is an antisense oligo used to elicit human RNase (ribonuclease) H degradation of target RNA. This sequence is used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 100.0%; Score 16; DB 1; Length 16; Similarity 100.0%; Pred. No. 1.1e+03; 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1434
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Gaps

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2709 AAAAAAAAAAAAA 2724

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16 AAAAAAAAAAAAA 1

6451/c AAD56451 standard; DNA; 16 BP.

RESULT 1433 AAD56451/ peptide nucleic acid; signalling;

hydroxyproline nucleic acid; HypNA; PNA; peptide nucleic acid gene expression; respiration; secretion; signalling; ion-channel activity; cell motility; developmental phenotype; tumour regression; hybridisation; ss.

DNA hybridisation oligomer SEQ ID 9.

(first entry)

ВР.

ADB68519 standard; DNA; 16

ADB68519;

/\*tag= a /note= "Optional N-terminal acetyl"

Location/Qualifiers

Key misc\_difference 1

Synthetic.

WO2003068798-A2 21-AUG-2003

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The invention relates to a novel assay for detecting a polynucleotide in a sample, which comprises: contacting a sample suspected of containing a carget polynucleotide with a predetermined single-estranded sensor polynucleotide complementary to the target polynucleotide, in a solution comprising an agent that is a nonaqueous solvent that allows the sensor comprising an agent that is a nonaqueous solvent that allows the sensor comprising an agent that is a nonaqueous solvent that allows the sensor polynucleotide, and determining fluorescence emission. The assay is useful for detecting a single or double-stranded target cassay is useful for detecting a single or double-stranded target comparison and polynucleotide, such as, DNA or RNA in a sample. The assay finds use in a polynucleotide, such as, DNA or RNA in a sample. The assay finds use in a corresic testing to identify the species or individual which was the source of a forensic specimen, in anthropological setting, paternity for eneming testing for compatibility between prospective tissue or blood conors and patients and in screening for hereditary disorders. The method is also useful to study alterations of gene expression in response to a stimulus, disease, drug or medication, and other applications include human haplotype diversity. The method is useful for detecting polynucleotide sequences from contaminants or pathogens including bacteria, yeast, and viruses to detect single nucleotide sequences from contaminants or pathogens including bacterial, yeast, and viruses to detect single nucleotide sequences from contaminants or pathogens of allelee. The method is useful for detection of mutations and to detect and alleles or the method is useful for detection of detections and to detect allowed and allelees of 
                                                                                                                                                                                                            Detection; single-stranded sensor; detectable fluorescence emission; forensic testing; paternity testing; tissue typing; hereditary disorder; human population genetics; human evolutionary history; cystic fibrosis; human haplotype diversity; Tay-Sachs; sickle-cell anaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polynucleotide sequence represents an oligonucleotide sequence used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorders including cystic fibrosis, Tay-Sachs, and sickle-cell anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting polynucleotides, for pharmacogenetic testing, comprises contacting a target polynucleotide with a complementary single-stranded sensor polynucleotide and an agent that allows the sensor to fluoresce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.6%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 1.1e+03; ve 0; Mismatches 0; Indels
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                                                                                                         Oligo-homodeoxyribonucleotide sequence, oligo dT.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L6-APR-2002; 2002WO-US012176.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-2001; 2001US-00836579.
06-MAR-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unidentified
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Method of inhibiting expression of genes or RNA transcripts, useful for therapy and determining effects of genes, by administering oligomers containing hydroxyproline nucleic acid.

Example 17; Page 233; 240pp; English.

Choob

Efimov V, Fernandez J, Archdeacon D, Archdeachon J,

(ACTI-) ACTIVE MOTIF

WPI; 2003-689653/65.

07-FEB-2003; 2003WO-US003904. 09-FEB-2002; 2002US-00072975.

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The invention relates to a novel method of inhibiting the expression of one or more genes or RNA transcripts by administering at least one oligonucleotide analogue that includes at least one hydroxyproline nucleic acid (HypNA) monomer to a cell or organism or their extracts. Throlling gene expression, respiration, secretion, signalling, ion-including gene expression, respiration, secretion, signalling, ion-expression. Furthermore, they may be utilised to determine the effects of particular genes, as antisense or homologous recombination constructs eg. for creating animal models of disease and finally, for increasing the activity of some enzymes, such as polymerases. The current sequence is that of the DNA hybridisation oligomer SEQ ID 9 of the invention. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence may also comprise a peptide nucleic acid (PNA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0;
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Best Local Similarity 100.0
Matches 16; Conservative
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Gaps

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AAAAAAAAAAAAA 1

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RESULT 1435 ADB68519/c

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31-MAY-2002; 2002US-0384454P.
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                                                              Kamme FC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEB77257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1438
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                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for nucleic acid analysis.

Specifically, it refers to a kit for performing nucleic acid detection using a chip to identify a target polynuclectide sequence. The present invention provides a chip (microarray) for nucleic-acid detection that significantly reduces non-specific binding produced between a test substance nucleic acid and a DNA probe. As such, this method can be used to detect the existence of a target sequence in a sample and for the analysis of polymorphisms, expression analyses, and presence or absence of a gene expression and SNPs, micro satellite sequences. Furthermore, of a gene expression and SNPs, micro satellite sequences. Furthermore, for diagnosis of disease by analyzing disease-related genes, estimation of incidence risk rate, detection of infected existence, analysis of virus type etc. This DNA microarray also has applications for various. Colinical objectives and for the inspection of food stuffs, quarantine, pharmaceutical, forensic medicine, agriculture, live stock forming, fishing and forestry. This oligonucleotide sequence is a used in the development of the DNA microarray of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                  complementary to target sequence and sequence existing in nature at low probability and probe for trapping which is complementary to medium
                        DNA detection; hybridization; DNA microarray; diagnosis; SNP detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                       Analyzing nucleic acid by use of medium probe consisting of sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid amplification; RNA transcription; RNA polymerase; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.6%; Score 16; DB 1; Length 16;
100.0%; Pred. No. 1.1e+03;
rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 16 BP; 16 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
 DNA oligo #1 related to the DNA microarray.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of an oligo dT16.
                                                                                                                                                                                                                                                                                                     Disclosure; Page 5; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2709 AAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI34487 standard; DNA; 16 BP
                                                                                                                                     26-DEC-2001; 2001JP-00395236.
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                                                                                                                                                              26-DEC-2001; 2001JP-00395236
                                     pharmaceutical; forensic; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 AAAAAAAAAAAAA 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16; Conservative
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                                                                                                                                                                                    (TOKE ) TOSHIBA KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003102243-A1.
                                                                                    JP2003189868-A.
                                                                                                             08-JUL-2003.
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                                                              Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   eliminating single-stranded oligonucleotide from the transcription sample. The method involves synthesizing single-stranded cDNA by incubating the sample RNA with reverse transcriptase and an oligonucleotide primer that primes synthesis in a direction toward 5' end of the RNA; converting the single-stranded cDNA into double-stranded cDNA to form a transcription sample containing a cDNA template; eliminating single-stranded oligonucleotide from the transcription sample; and transcribing the cDNA template into RNA using an RNA polymerase. The method is useful for improving RNA polymerase based RNA transcription template derived production of RNA in the transcriptis the undesired nontemplate derived production of RNA in the transcription reaction. Sequences AD134483-AD134489 represent oligonucleotides used in a 77 RNA
                                                                                                                                                                                                             Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo, SEQ ID NO: 1, to make full-length coding sequence cDNA libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Making full-length coding sequence cDNA libraries for research purposes comprises binding a tag molecule (e.g. biotin or avidin) to a diol structure present in the 5' cap site of an mRNA forming an RNA-DNA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to amplifying for RNA in a sample comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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Pred. No. 1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 6; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.6%; Scc-
100.0%; Prev
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Best Local Similarity 100.
Matches 16; Conservative
(JANC ) JANSSEN PHARM NV.
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                                                                                                                                           WPI; 2004-035466/03.
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                                                                     Zhu JY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA library; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2005175993-A1
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The present invention relates to a method for making cDNA libraries wherein the CDNA inserts comprise the full-length of the mcding sequences but but having lengths less than the full-length of the mRNA. The method of the invention comprises binding a tag molecule to a diol structure present in the 5' Cap sites of mRNAs, forming RNA-DNA hybrids in to synthesize the first cDNA strand, spearating RNA-DNA hybrids carrying a DNA corresponding to full-length of mRNAs from RNA-DNA hybrids formed above by using a function of the tag molecule and strand, aperating the second cDNA strand by self-priming the first cDNA strand by contain the full-length of the mRNAs but do contain the full-length of the coding sequences of the mRNAs. The invention is useful for making or constructing full-length coding sequence cDNA libraries which may be utilized in researches in the fields of medical science and biology. The present sequence is an in the fields of medical science and biology. The present sequences cDNA
                          Disclosure; SEQ ID NO 1; 13pp; English.
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0.6%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 1.1e+03; ive 0; Mismatches 0; Indels Sequence 16 BP; 16 A; 0 C; 0 G; 0 T; 0 U; 0 Other; 2709 AAAAAAAAAAAAA 2724 Query Match 0.6 Best Local Similarity 100. Matches 16; Conservative ò

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Gaps ö

> 1 AAAAAAAAAAAAA 16 엄

drought resistance; crop improvement; ZmRSPTP1; ss. Zea mays ZmRSPTP1 associated oligonucleotide #9. AEC34066 standard; DNA; 16 BP 03-NOV-2005 (first entry) Unidentified CN1584034-A. AEC34066; RESULT 1439 

21-AUG-2003; 2003CN-00153941 21-AUG-2003; 2003CN-00153941 23-FEB-2005

(BEIJ-) BEIJING AGRIC BIOTECHNOLOGY RES CENT

Huang C; Wu Z, Jia W,

WPI; 2005-406151/42.

Corn tyrosin protein phosphatase gene and its coding protein and use. Example 2; Page 6; 13pp; Chinese. The invention describes a ZmRSFTP1 gene, its coding protein and use. The EmRSFTP1 gene is one of the following ribonucleotide sequence: 1) SEQ ID No:1 in sequential table; 2) ribonucleotide sequence of coded SEQ ID No:2 protein sequence; 3) DNA sequence having 90% homology with DNA sequence refined by SEQ ID No:1 in sequentialtable. It can be used to breed drought-resistant plant and corn. This sequence represents an oligonucleotide associated with the isolation of ZmRSPTP1.

Seguence 16 BP; 1 A; 0 C; 0 G; 15 T; 0 U; 0 Other;

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           Gaps
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0.6%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 1.1e+03;
                                                                                                                gene amplification; gene mapping; gene sequencing; ss.
                                                                                                    Family 16/15{inv(alpha.alpha)]:4 nucleotide fragment.
           Mismatches
                                                                                                                                                                               /mod_base= other
/note= "3'-phosphorylated"
                                                                                                                                                             'note= "5'-phosphorylated"
                                                                                                                                      Location/Qualifiers
0.00,
100.0%; Pre
                                                                                                                                                   /*tag= a
/mod base= other
                        2708 TAAAAAAAAAAAA 2723
                                                                BP.
                                                                AED63168 standard; DNA; 16
                                                                                       (first entry)
                                   16 TAAAAAAAAAAAAA
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            16; Conservative
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Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                                                            modified base
                                                                                                                                                                    modified_base
                                                                                        29-DEC-2005
                                                                                                                           Synthetic.
                                                                             AED63168;
                                                      RESULT 1440
                                                           AED63168,
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08-APR-2005; 2005WO-US011812. 09-APR-2004; 2004US-0563283P. 26-APR-2004; 2004US-0565284P. (UYBO-) UNIV BOSTON 27-OCT-2005

Siddigi FA;

Cantor CR,

WPI; 2005-725959/74.

Determining a target nucleic sequence comprises cleaving transcript of sequence in a sequence-specific manner, determining molecular weight of fragments, performing fragment identity mapping and comparing the observed mass.

Example 1; SEQ ID NO 27; 92pp; English.

The invention relates to determining a target sequence of a template

nucleic acid. The method involves: creating a transcript of an isolated

cuplete nucleic acid using polymerase enzyme and nucleosides selected

corresponding to the sequence specific reactions resulting in complete cleavage of

primers; performing a cleavage reaction resulting in complete cleavage of

the transcript in a sequence specific manner into fragments using cutters

selected from the group consisting of enzymmaic cutters, chemical

cutters, and their combination, analyzing the cleavage reaction products

cutters and their combination; analyzing the cleavage reaction products

cutters and the molecular weights of the fragments performing fragment

identity mapping using nucleotide masses and cleavage reactions,

che cutters to calculate the molecular weights and sequences of all

consible fragments that result from the second step cleavage reactions,

and comparing the masses observed in the third step with the fragment

cleantity mapping, where the comparison results in determination of all

cleantity mapping, where the comparison results in determination of all

che target sequences present in the sample. In determination of all

sequence of a template nucleic acid, the steps are performed at least two

times with different cutters, thus allowing production of overlapping

fragments, and compiling the overlapping fragments to produce at least

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to determining a target sequence of a template nucleic acid. The method involves: creating a transcript of an isolated template nucleic acid using polymerase enryme and nucleosides selected for sequence specific reactivity and molecular weight and oligonucleotide primers; performing a cleavage reaction resulting in complete cleavage of
one larger subsequence. The larger subsequence is the complete sequence of the template. The primers are sequence specific and have a random sequence. The molecular weight is determined using mass spectroscopy which is matrix-assisted laser desorption/ionization time-of-flight spectroscopy. The method allows de novo detection of sequences in a target nucleic acid without requiring any prior sequence information. Also provided is a method for determining the number of genes in a nucleic acid sample. Sequences AED63145-AED63168 represent modified fragments produced during fragment identity mappings for 16/15[inv(alpha.alpha)]:4 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining a target nucleic sequence comprises cleaving transcript of sequence in a sequence-specific manner, determining molecular weight of fragments, performing fragment identity mapping and comparing the observed mass.
                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                   0.6%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 1.1e+03; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene amplification; gene mapping; gene sequencing; ss.
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                                                                                                                                                                                    Sequence 16 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
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/note= "3'-phosphorylated"
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mod base= other
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26-APR-2004; 2004US-0565284P.
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the transcript in a sequence-specific manner into fragments using cutters selected from the group consisting of enzymatic cutters, chemical cutters, and their combination; analyzing the clasvage reaction products to determine the molecular weights of the fragments; performing fragment identity mapping using nucleotide masses and cleavage specificities of the cutters to calculate the molecular weights and sequences of all possible fragments that result from the second step cleavage reactions; and comparing the masses observed in the third step with the fragment of and comparing the masses observed in the third step with the fragment of the target sequences present in the sample. In determining a target sequence of a template nucleic acid, the steps are performed at least two times with different cutters, thus allowing production of overlapping fragments and compiling the overlapping fragments to produce at least conclusions and surpayers subsequence is the complete sequence of the template. The primers are sequence specific and have a random sequence. The melecular weight is determined using mass spectroscopy which is matrix-assisted laser desorption/ionization time-of-flight contraction of sequences in a contract of carry in the nucleic acid without requiring any prior sequence in a new product in a contract in the number of cannes in a new part of cannes in a new product of cannes in a new part of 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Also provided is a method for determining the number of genes in a nucleic acid sample. Sequences AED63145-AED63168 represent modified fragments produced during fragment identity mappings for 16/15[inv(alpha.alpha)]:4 family.
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Matches 16, Conservative
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of nucleic acid molecules from the present invention
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                                                     Query Match
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                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fims-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule modulating VEGF receptor(s) gene expression or mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0;
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  Claim 4; Page 79; 218pp; English.
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Best Local Similarity 100.
Matches 16, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA and encoded protein which activates eosinophil cells - for treatment of cancer, parasite infection, autoimmune disease and allergic inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bosinophil cell activator; treatment; diagnosis; malignant tumour;
parasitic infection; allergic inflammation; eosinophilic pneumonia;
rapid onset eosinophilia; autoimmune disease; gene therapy; primer; 88.
                                                                                                        Gaps
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Pred. No. 1.1e+03;
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Sequence 17 BP; 0 A; 1 C; 0 G; 0 T; 16 U; 0 Other;
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100.0%; Pred. No. 1.1e+03;
rative 0; Mismatches 0:
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tive 0; Mismatches
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nes 16; Conserv
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06-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-APR-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Obayashi I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                       sednences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nagasu T,
                                                                                                                                                                                                                                                                                                                                                                                                                               AAA30179;
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1447
                                                                                                                                                                                                                                                                                                                                                                                               AAA30179,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                              This sequence represents a primer of the invention. The invention relates to sequences of at least two nucleotides of formula: (X)m5'-(alpha)n-beta and; or (X)m5'-(almma)k-delta-N3'; where X = a labelled compound and/or nucleotide with voluntary sequence; m = 0 or 1; alpha = thymine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, guanine or cytosine; N = adenine, guanine or thymine; and = thymine; a natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, guanine and/or cytosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                         Peptides having at least two new nucleotides - useful as primers in RT-
                                                                           RT-PCR primer; DNA sequence determination; gene sequence analysis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RT-PCR primer; DNA sequence determination; gene sequence analysis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16; DB 1; Le
Pred. No. 1.1e+03;
0; Mismatches 0;
                                                      RT-PCR primer of the invention SEQ ID 12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RT-PCR primer of the invention SEQ ID 11.
                                                                                                                                                                                                                                                                                         Disclosure; Page 11; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.6%; Scc...
100.0%; Pre
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                                                                                                                                                                   97JP-00208312.
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                                (first entry)
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                                                                                                                                                                                                                                     WPI; 1999-183822/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                 11-MAY-1999
                                                                                                                      JP11032765-A
                                                                                                                                                                                        18-JUL-1997;
                                                                                                                                                                 18-JUL-1997;
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                                                                                                                                            09-FEB-1999,
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                                                                                                  Synthetic
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           AAX18371;
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This sequence represents a primer of the invention. The invention relates to sequences of at least two nuclectides of formula: (X)m5'-(alpha)n-beta -N3'; or (X)m5'-(gamma) **Ledelta-N3'; where X = a labelled compound and/or a nucleotide with voluntary sequence; m = 0 or 1; alpha = thywine; n = natural number indicating the repetition of alpha; beta, delta = V or N; V = adenine, quanine or cytosine; N = adenine, guanine, cytosine or thymine; gamma = thymine; k = natural number of 3 or over indicating the repetition of gamma, in which thymine expressed by gamma is composed of 1/3 or less of adenine, quanine and/or cyrosine. The new nucleotides are useful as primers for RT-PCR and determination of base sequences. The new sequences allow for reproductive and highly efficient analysis of gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This sequence represents a PCR primer used in the identification of a human pollenosis associated gene. The gene is highly expressed in individuals with high pollen-specific immunoglobulin E (IgE) levels. The
                                                                                  Peptides having at least two new nucleotides - useful as primers in RT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gunji S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gene highly expressed in patients with high cedar pollen-specific IgE levels, useful for diagnosing pollenosis, and screening candidate compounds for pollenosis treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pollenosis-associated protein; high pollen-specific immunoglobulin E; 19E; diagnose; cedar pollenosis; treatment; human; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer GT15A used in pollenosis associated gene identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16; DB 1; Length 17;
Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 2 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oshida T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.6%; Scor.
100.0%; Pred. No. ...
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Lu N, Ogawa K;
                                                                                                                                                   Disclosure, Page 11, 19pp, Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Page 38; 44pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2708 TAAAAAAAAAAAA 2723
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(TAKI ) TAKARA SHUZO CO LTD.
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Best Local Similarity 100.
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sugita Y,
Imai Y,
                                          WPI; 1999-183822/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-317712/27.
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invention relates to the nucleotide sequence encoding the pollenosis associated protein, diagnosing pollenosis and screening candidate compounds for treating pollenosis. The gene can be used in diagnosing pollenosis, particularly cedar pollenosis, and screening candidate compounds for pollenosis treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kuga T;
                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (gA nephropathy-associated protein; diagnosis; treatment; antisense;
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Sawada S, Takei M, Shibata K, Furuya A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA sequences preferentially expressed in IgA nephropathy proteins encoded by them, and antibodies to those proteins
                                                                                                                                                        DB 1; Length 17; 1.1e+03;
                                                                                                                                                                                             0; Indels
                                                                                                                    Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human IgA nephropathy-associated cDNA primer #61.
                                                                                                                                                        0.6%; Score 16; DB 100.0%; Pred. No. 1.1 Live 0; Mismatches
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                                                                                                                                                                                                                                       2708 TAAAAAAAAAAAAA 2723
                                                                                                                                                                                                                                                                                                                                                                          AAX82720 standard, DNA; 17 BP.
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                                                                                                                                                                                                                                                                           17 TAAAAAAAAAAAA 2
                                                                                                                                                    Query Match 0.6%
Best Local Similarity 100.0°
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                     RESULT 1448
                                                                                                                                                                                                                                                                                                                                                         AAX82720,
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display, in the method of the invention. The specification describes a method for measuring the level of two or more nucleic acid molecules a method for measuring the level of two or more nucleic acid molecules in a target. The method comprises contacting a probe with an arbitrarily or statistically sampled target and detecting the amount of specific binding of the farget to the probe. The methods can be used to identify contacting the amount of specific binding differentially expressed nucleic acid molecules associated with disease states, such as cancer, autoimmune disease, infectious disease, aging, a developmental disorder, proliferative disorder or neurological disorder. Alternatively the methods can be used to assess the efficacy or toxicity of or a resistance to a treatment. Also the methods can be used to determine differential expression of nucleic acid molecules in response to a stimulus, e.g. a chemical, drug or growth factor (especially especially can also be used to determine co-regulated genes that can be potential
                                                                                                                                                                                                                                                               Stimulus-regulated nucleic acid; sequence profile; nucleic acid level; differentially expressed nucleic acid; disease state; cancer; autoimmune disease; infectious disease; aging; developmental disorder; proliferative disorder; neurological disorder; toxicity; primer; treatment resistance; differential expression; drug discovery; growth factor; epidermal growth factor; radiation; stress; pathogen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Measuring expression of low abundance reduced complexity target nucleic
                                                                                                                                                                                                                                  Anchored oligo(dT) primer AT15A used for modified differential display
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 91; 187pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KIMM-) KIMMEL CANCER CENT SIDNEY.
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tive 0;
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                                                                                                                           AAZ36739 standard; DNA; 17
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Best Local Similarity 100.0
Warches 16; Conservative
17 TAAAAAAAAAAAAA
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                                                                                         RESULT 1449
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Query Match 0.6%; Score 16; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 1.1e+03; Matches 16; Conservative 0; Mismatches 0; Indels

Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;

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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclase activity. (A), and moor generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium).

Complete acid (A') that modulates expression of the oestrogen receptor in vivo or by transforming cells ex vivo and implanting treated cells, or for other conditions associated with levels of oestrogen receptor.

Correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of therapeutic targets, and as research ceststance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences.

CAAA2474 represent oestrogen receptor hammerhead ribozyme sequences.

CAAA2474 represent oestrogen receptor hairpin ribozyme sequences.

CR AAA2474 represent oestrogen receptor hairpin ribozyme sequences.

CR CAAA26105 represent oestrogen receptor hairpin ribozyme sequences.

CR CAAA26105 represent oestrogen receptor hairpin ribozyme sequences.

CR CAAA26105 represent oestrogen receptor hairpin ribozyme and antisense oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids that interact, and optionally cleave, target sequences,
                                                                                                                                                                   Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1947.
                                                                                                                                                                                                       Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense Oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bellon L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Karpeisky A,
Haeberli P;
                                                                                                                                                                                                                                                                           anticancer; breast cancer; endometrium cancer; ss.
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Reynolds M, Zwick M, Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 77; Page 79; 148pp; English.
                                          AAA25449 standard; DNA; 17 BP.
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98US-00103636.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC
                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used to treat cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matulic-Adamic J;
                                                                                                                              19-JUL-2000
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                                                                                    AAA25449;
RESULT 1450
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0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03;
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100.0%; Pro 2709 AAAAAAAAAAAAA 2724 Query Match 0.69 Best Local Similarity 100.0 Matches 16; Conservative

ò g

17 AAAAAAAAAAAAA 2

RESULT 1451 AAA25451/c

AAA25451 standard; DNA; 17 BP.

19-JUL-2000

(first entry)

Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1949. Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.

Homo sapiens

WO9954459-A2.

28-OCT-1999.

99WO-US008547. 19-APR-1999;

98US-00103636. 98US-0082404P. 20-APR-1998; 23-JUN-1998;

(RIBO-) RIBOZYME PHARM INC.

Bellon L; Karpeisky A, Haeberli P; Beigelman L, Mcswiggen JA, Zwick M, Jarvis T, Woolf T, Matulic-Adamic J; Thompson JD, Reynolds M,

WPI; 2000-013248/01.

New nucleic acids that interact, and optionally cleave, target sequences, used to treat cancer.

Claim 77; Page 79; 148pp; English.

The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate (Ink, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium). (C) in vivo or by transforming cells as vivo and implanting treated cells, or for other conditions associated with levels of estrogen receptor. (C) for other conditions associated with levels of estrogen receptor. (C) Escause of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research creagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves consistence to nucleases, binding affinity and/or activity. AAA23503 to AAA23593 to AAA25010 represent cestrogen receptor hairpin ribozyme sequences. (C) AAA24747 represent cestrogen receptor hairpin ribozyme sequences. (C) AAA25010 to AAA2501 invention

Sequence 17 BP; 0 A; 0 C; 1 G; 16 T; 0 U; 0 Other;

ö 0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Indels 16; Conservative Local Similarity Query Match Matches

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Gaps

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Gaps

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0; Indels

Mismatches

RESULT 1452 AAC64202/c ID AAC6420: XX

AAC64202 standard; DNA; 17 BP.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to the human pollinosis-associated gene 373 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates also relates to the protein encoded by pollinosis associated gene 373; expression constructs and host cells comprising pollinosis-associated gene 373 nucleic acids; pollinosis-associated gene 373 nucleic acids; and a method of pollinosis-associated gene 373 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 373 nucleic acids. The invention additionally encompasses methods of screening drug acids. The invention additionally encompasses methods of screening drug acids. The invention additionally encompasses methods of screening drug expression of pollinosis-associated gene 373 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control.

Pollinosis-associated gene 373 is useful in the diagnosis of allergic diseases and in the screening of drug candidates for the treatment of allergic diseases to the treatment of sequence represents a PCR primer used in the greening of drug candidates for the treatment of sequence represents a PCR primer used in the
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                                                                                                                                                                                                                                                                                                                                                                                        Pollinosis-associated gene 373 undergoing significantly low expression in subjects with high cedar pollen-specific immunoglobulin-E levels, useful in diagnosis of allergic diseases and screening drug candidates.
                                                                                  Human; pollinosis-associated gene 373; IgB; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; diagnosis; drug screening; allergic disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                     Gunji S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                         PCR anchor primer, SEQ ID NO:3, used in human gene 373 isolation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR anchor primer, SEQ ID NO:2, used in human gene 419 isolation.
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                                                                                                                                                                                                                                                                                                                     Obayashi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.6%; Score 16; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 1.1e+03; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             isolation of human pollinosis-associated gene 373 cDNA
                                                                                                                                                                                                                                                                                                                   Kashiwabara T, Oshida T, Obay
Yoshida N, Ogawa K, Matsui K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 6; Page 69; 80pp; Japanese.
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                                                                                                                                                                                                                                  26-APR-2000; 2000WO-JP002730.
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                              (first entry)
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                                                                                                                                                                                                                                                                                        (GENO-) GENOX RES INC.
                                                                                                                                                                                                                                                                                                                     Sugita Y,
, Imai Y,
                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-687339/67.
                                                                                                                                                                         WO200065046-A1
                                                                                                                                                                                                                                                               27-APR-1999;
                                                                                                                                                                                                                                                                                                                     Nagasu T, Si
Obayashi I,
                               21-FEB-2001
                                                                                                                                                                                                      02-NOV-2000
                                                                                                                                              Synthetic.
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   AAC64202
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The invention relates to the human pollinosis-associated gene 419 which exhibits reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 419 has homology with the gene encoding human Fas-associated factor-1 (FAF-1). The invention also constructs and host cells comprising pollinosis-associated gene 419; expression constructs and host cells comprising pollinosis-associated gene 419 primers and probes; antibodies against the protein encoded by the gene; methods of detection of pollinosis-associated gene 419 uncleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 419 uncleic acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic diseases via the presence of a test compound relative to a control.Pollinosis-associated gene 419 in the presence of a test compound relative to a control.Pollinosis-associated gene 419 in the diagnosis of allergic diseases and in the screening of drug candidates (for the treatment of squed diagnosis of allergic diseases. The present sequence represents a PCR for the treatment of squed diagnosis of allergic diseases. The present sequence represents a PCR for the treatment of squed diagnosis of allergic diseases. The presents as a PCR for the treatment of squed diseases. The presents appresents a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pollinosis-associated gene 419 undergoing significantly low expression in subjects with high cedar pollen-specific IgE levels, useful in diagnosis of allergic diseases and screening drug candidates.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gunji S;
Human, pollinosis-associated gene 419; FAF-1 homologue;
Fas-associated factor-1; IgE; immunoglobulin E; cedar pollen allergy;
T-cell; reduced expression; detection; diagnosis; drug screening;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sugita Y, Kashiwabara T, Oshida T, Obayashi M,
Imai Y, Yoshida N, Ogawa K, Matsui K;
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100.0%; Pred. No. 1.1e+03;
[ve 0; Mismatches 0;
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100.0%; Pre-
                                                                                                                           allergic disease; PCR primer; ss.
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GENO-) GENOX RES INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nagasu T, Sı
Obayashi I,
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                                                                                                                                                                                                                     Synthetic.
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The invention relates to the human pollinosis-associated gene 513 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates to methods of detection of pollinosis-associated gene 513 nucleic acids; a method of diagnosis of allergic diseases via the detection of pollinosis-associated gene 513 nucleic acids; and methods of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 513 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 513 in the diagnosis of allergic diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-
                                                                                                                                                                                                                                                                                                           Pollinosis-associated gene 513 undergoing significantly low expression in subjects with high cedar pollen-specific IgE levels, useful in diagnosis of allergic diseases and screening drug candidates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, pollinosis-associated gene 581, IgE; immunoglobulin E, cedar pollen allergy; T-cell; reduced expression, detection; diagnosis; drug screening; allergic disease; PCR primer; ss.
                                                                                                                                                                                                                                  Gunji S;
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                                                                                                                                                                                                                                Obayashi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 17;
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                                                                                                                                                                                                                               T, Oshida T, Obay
Ogawa K, Matsui K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
drug screening; allergic disease; PCR primer; ss.
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Pred. No. 1.1e+03;
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                                                                                                                                                                                                                             Sugita Y, Kashiwabara T,
Imai Y, Yoshida N, Oga
                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 38; 46pp; Japanese.
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100.0%; Pre
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                                                                                                                              26-APR-2000; 2000WO-JP002733.
                                                                                                                                                              99JP-00120491
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                                                                                                                                                                                              (GENO-) GENOX RES INC.
                                                                                                                                                                                                                                                                             WPI; 2000-687342/67.
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                                                               WO200065049-A1
                                                                                                                                                                27-APR-1999;
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Obayashi I,
                                                                                               02-NOV-2000
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                                  Synthetic.
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The invention relates to the human pollinosis-associated gene 581 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific Ig8 (immunoglobulin B) levels. The gene cwas isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates also relates to the protein encoded by pollinosis-associated gene 581; to expression constructs and host cells comprising pollinosis-associated gene 581 mucleic acids; pollinosis-associated gene 581 primers and probes; antibodies against the protein encoded by the gene; methods of detection of pollinosis-associated gene 581 nucleic acids; and a method of diagnosis of allergic diseases via the detection of pollinosis-associated companied of gene 581 nucleic acids; and a method of companied to a pollinosis-associated gene 581 nucleic acids; and a method of companied to a control acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic disease by control. Pollinosis-associated gene 581 is useful in the diagnosis of a control. Pollinosis-associated gene 581 is useful in the diagnosis of allergic diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer control in the inclain of human nollinosis-associated gene 581 cDNA expression in in diagnosis Human, pollinosis-associated gene 627, IgE; immunoglobulin E; cedar pollen allergy; T-cell; reduced expression; detection; diagnosis; drug screening; allergic disease; PCR primer; ss. Gunji S; Gaps used in the isolation of human pollinosis-associated gene 581 cDNA PCR anchor primer, SEQ ID NO:2, used in human gene 627 isolation. ; Obayashi M, Pollenosis-associated gene 581 undergoing significantly low subjects with high cedar pollen-specific IgE levels, useful of allergic diseases and screening drug candidates. Score 16; DB 1; Length 17; Pred. No. 1.1e+03; 0; Indels Kashiwabara T, Oshida T, Obay Yoshida N, Ogawa K, Matsui K; Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other; Query Match 0.6%; Score 16; UB Best Local Similarity 100.0%; Pred. No. 1.1 Example 6; Page 39; 69pp; Japanese 2708 TAAAAAAAAAAAA 2723 BP 99JP-00120492 26-APR-2000; 2000WO-JP002735. 99JP-00120493. 17 TAAAAAAAAAAAAA 2 AAC64213 standard; DNA; 17 (first entry) (GENO-) GENOX RES INC. Sugita Y, Imai Y, (GENO-) GENOX RES INC WPI; 2000-687341/67. WO200065051-A1 27-APR-1999; Nagasu T, S Obayashi I, 27-APR-1999; 21-FEB-2001 02-NOV-2000. Synthetic. AAC64213; RESULT 1456 AAC64213, 8 셤 

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The invention relates to the human pollinosis-associated gene 627 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IGE (immunoglobulin B) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. The invention also relates to methods of detection of pollinosis-associated gene 627 nucleic acids; amethod of diagnosis of allergic diseases via the detection of pollinosis-associated controlled of allergic diseases by measuring the expression of pollinosis-associated controlled and a method of screening drug candidates for the associated gene 627 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 627 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 627 in pollen diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-Pollinosis-associated gene 627 undergoing significantly low expression in subjects with high cedar pollen-specific IgE levels, useful in diagnosis of allergic diseases and screening drug candidates. ŝ Obayashi M, Gunji Kashiwabara T, Oshida T, Obay: Yoshida N, Ogawa K, Matsui K; Example 6; Page 41; 51pp; Japanese Sugita Y, Imai Y, WPI; 2000-687344/67 Obayashi I, Nagasu T,

Seguence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;

Gaps . 0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Indels Query Match Best Local Similarity 100. Matches 16; Conservative

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AAC64230 standard; DNA; 17 BP AAC64230; RESULT 1457 PRACE TO THE PRACE OF THE PRACE

(first entry) 21-FEB-2001 PCR anchor primer, SEQ ID NO:2, used in human gene 795 isolation.

Human; pollinosis-associated gene 795; vimentin homologue; IgB; immunoglobulin B; cedar pollen allergy; T-cell; reduced expression; detection; diagnosis; drug screening; allergic disease; PCR primer; ss.

Synthetic.

WO200065050-A1.

02-NOV-2000

26-APR-2000; 2000WO-JP002734

99JP-00120494 27-APR-1999;

(GENO-) GENOX RES INC. (EISA ) EISAI CO LTD.

Sugita Y, Imai Y,

Nagasu T, S Obayashi I,

Yokoi A;

Pollinosis-associated gene 795 undergoing significantly low expression in subjects with high cedar pollen-specific IgE levels, useful in diagnosis WPI; 2000-687343/67.

of allergic diseases and screening drug candidates.

Page 45; Example 6; 73pp; Japanese.

The invention relates to the human pollinosis-associated gene 795 which exhibits significantly reduced expression in the T-cells of individuals with high cedar pollen-specific IgE (immunoglobulin E) levels. The gene was isolated from T-cells from individuals allergic to cedar pollen using the differential display method. Pollinosis-associated gene 795 has concerned to the protein encoded by pollinosis gene 795; to expression relates to the protein encoded by pollinosis gene 795; to expression contains the protein encoded by pollinosis associated gene 795 constructs and host cells comprising pollinosis-associated gene 795 nucleic acids, pollinosis expression of pollinosis of allergic diseases via the detection of pollinosis-associated gene 795 nucleic acids. The invention additionally encompasses methods of screening drug candidates for the treatment of allergic disease by constructing the expression of pollinosis-associated gene 795 in pollen and an expression of pollinosis-associated gene 795 in pollen a control Pollinosis-associated gene 795 in pollen a control Pollinosis-associated gene 795 in pollen a control pollinosis-associated gene 795 in pollen treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-associated gene 795 cDNA a control. Pollinosis-associated gene 795 is useful in the diagnos allergic diseases and in the screening of drug candidates for the treatment of such diseases. The present sequence represents a PCR 

Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;

Gaps .. 0 0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Indels Best Local Similarity 100. Matches 16; Conservative Query Match

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BP. AAC92292 standard; DNA; 17 (first entry) 22-MAR-2001 AAC92292; RESULT 1458 AAC92292 

Human pollinosis-associated gene 465 related PCR primer SEQ ID NO:2.

Human; pollinosis-associated gene 465; pollen scattering; allergy; allergic disease; PCR primer; ss.

Homo sapiens.

WO200073439-A1.

07-DEC-2000

18-MAY-2000; 2000WO-JP003191.

99JP-00148784 (GENO-) GENOX RES INC. (EISA ) EISAI CO LTD. 27-MAY-1999;

Nagasu T, Sugita Y, Kashiwabara T, Oshida T, Obayashi M, Gunji Obayashi I, Imai Y, Yoshida N, Ogawa K, Matsui K, Takahashi E; Yokoi A;

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WPI; 2001-061528/07.

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Kashiwabara T, Oshida T, Obayashi M, Gunji Yoshida N, Ogawa K, Matsui K, Takahashi E;

Pollinosis-associated gene 465 undergoing significantly low expression in subjects after pollen scattering, useful in diagnosis of allergic diseases and screening candidate compounds to regulate response of T cells to antigen stimulus.

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                                                               The present invention describes the human pollinosis-associated gene 465 which has a nucleic acid sequence of 3442 base pairs (bp), given in (AAC92291), that undergoes significantly low expression in subjects after pollen scattering, and is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The gene is useful in the diagnosis of allergic diseases and screening candidate compounds for remedies capable of regulating the response of T cells to the stimulus by an antigen. The present sequence represents a PCR primer which is used in an example from the present invention
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Yoshida N, Ogawa K, Matsui K, Takahashi
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reduced expression; detection; diagnosis; drug screening;
allergic disease; PCR primer; ss.
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Pred. No. 1.1e+03;
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Local Similarity 100.0%; Pred. No.
Page 43; 61pp; Japanese.
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, Imai Y,
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additionally encompasses a method of screening drug candidates for the treatment of allergic disease by measuring the expression of pollinosis-associated gene 787 in pollen antigen-stimulated T-cells in the presence of a test compound relative to a control. Pollinosis-associated gene 787 is useful in the diagnosis of allergic diseases and in the screening of argregation of the treatment of such diseases. The present sequence represents a PCR primer used in the isolation of human pollinosis-associated gene 787 cDNA
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Pred. No. 1.1e+03;
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pollen scattering; antigen; primer; ss.
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wa K, Matsui K;
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0;
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Yoshida N, Ogawa K,
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100.0%; Pred
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Best Local Similarity 100.
Matches 16; Conservative
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Imai Y,
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Bauren G;

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The present invention relates to a method for providing a profile of mRNA molecules present in a sample. The method comprises generating two independent patterns characteristic of the population of mRNA molecules expressed in the sample and analysing the patterns using a combinatorial algorithm, comparing gene expression by different or same cell types under different conditions, and identifying genes having a role in various cellular processes. The method is useful for the analysis and identification of transcribed genes, and fingerprinting. The method can be used to identify genes which play a role in determining various cellular processes, including susceptibility to external factors, development, and disease. The present sequence for a PCR primer is used in the production of a single pattern characteristic of a sample, employing a Type IIS restriction enzyme (i.e. FoXI) in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human, ss, PCR, acetyltransferase, 20-90-05; allergic disease, primer, differential display, eosinophil; antiallergic; atopic dermatitis; GT15A.
                                                                                                                                                                Providing mRNA profile, by generating two independent patterns characteristic of sample mRNA population, analyzing patterns, comparing gene expression by cell types under varied conditions, and identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Acetyltransferase-like protein 20-90-05 PCR primer GT15A.
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(NIGE-) JAPAN GEN NAT CHILDREN'S HOSPITAL.
(BISA ) EISAI CO LTD.
                                                                                                                                                                                                                                           Disclosure; Fig 2; 67pp; English.
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23-JUL-2001; 2001WO-IB001539.
                               21-JUL-2000; 2000GB-00018016.
21-JUL-2000; 2000US-0219925P.
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                                                                                                          Linnarsson S, Ernfors P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 100.
nes 16; Conservative
                                                                           (GLOB-) GLOBAL GENOMICS
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                                                                                                                                      WPI; 2002-217065/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention
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Takahashi E;
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides a method of diagnosis of allergies that involves: assaying the levels of expression of genes B101, B1466, B1072 or B1151 in T-cells; and comparing them with the level of expression in healthy T-cells. The method is useful for diagnosing allergies, particularly atopic dermatitis. The present sequence represents a PCR primer used for analysis of the expression of the above genes
                                                                                                                                                                    B1466; B1072; B1151; T-cell; allergy; atopic dermatitis; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                Saito
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diagnosis of allergies including atopic dermatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                            Matsui K,
                                                                                                                                                                                                                                                                                                                                                                 GENOX RES INC.
JAPAN GEN NAT CHILDREN'S HOSPITAL.
                                                                                                                                       Nucleotide sequence of primer GT15A.
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                                                                                                                                                                                                                                                                                                                                                                                                            Obayashi I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2708 TAAAAAAAAAAAAA 2723
                                              AAH47126 standard; DNA; 17 BP
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Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-557789/62
                                                                                                                                                                                  PCR primer; ss
                                                                                                                                                                                                                                               WO200165259-A1
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                  RESULT 1461
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Score 16; DB 1; Length 17; Pred. No. 1.1e+03; 0; Mismatches 0; Indels

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Tsujimoto G;

Nagasu T,

Gaps

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WPI; 2002-439993/47.
                                                                    Local Similarity
                                                                                                                 JP2002095500-A.
                                                                                                             Homo sapiens
                                                                      16;
                                                                                                                     02-APR-2002
                                                                              17
                                                                                            ABL59038;
                                                                  Query Match
                                                        display
                                                                                                                                 (GENO-)
(KOKU-)
                                                                                     1464
                                                                      Matches
                                                                                      ABL59038/
8
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The specification describes a method for examining allergosis. The method comprises measuring the expression level of the gene given in ABL59037, and comparing it with the expression level of the gene in the eosinophils of a healthy person. The method is used for the examination of allergosis. The present sequence represents a PCR primer, which is used in the course of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                which involves determining the expression level of a gene, having one of the 17 nucleotide sequences shown in ABN99812-ABN99828, in the eosinophils in a patient and comparing the expression level with that in the eosinophils of a healthy individual. The method can be used to examine allergic diseases, particularly atopic dermatitis, and its early diagnosis, which is also applicable in screening candidate compounds for remedies. The present sequence is a PCR primer described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Method for examining allergic diseases by differential display of seventeen genes showing different expression particularly significant increase in eosinophils in patients with mild atopic dermatitis, also applicable in screening compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; allergy; atopic dermatitis; eosinophil; anti-allergic; PCR;
                                                                                                                                                                                                                                                                                  0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; ive 0; Mismatches 0; Indels
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human allergic disease related PCR primer SEQ ID NO: 18.
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                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
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(NIGE-) JAPAN GEN NAT CHILDREN'S HOSPITAL.
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Example 1; Page 17; 20pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                   2708 TAAAAAAAAAAAA 2723
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Best Local Similarity 100.
Matches 16; Conservative
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Matches 16; Conservative
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8$88888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method for examining allergic diseases comprises determining the expression level of a gene containing, the human cDNA appearing as ABK49633 which has homology with a cetyltransferases in the eosinophils of a patient and comparing the expression level with that in the eosinophils of a healthy individual compounds which affect the expression level of the gene or the candidate compounds which affect the expression level of the gene or the activity of the protein encoded by the gene (including related proteins and mutants), the use of probes based on the gene sequence in the carmination of allergic diseases, the use of reporter constructs in the correction of allergic diseases, the use of reporter constructs in the correction of an endiance of the gene, cells transformed with the vector, an antibody against the protein and a model animal for allergic diseases (which is a transgenic non-human vertebrate with lowering of expression intensity of the gene in eosinophils. The method is examining allergic diseases particularly atopic dermattits which is also applicable in screening candidate compounds for remedies. Such method can be performed in high throughput, at low cost. The present sequence is a differential
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                                                                                                      stage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              high throughput, at low cost. The present sequence is a differential splay PCR primer for the cDNA encoding the human acetyltransferase-like
                                                             Examining allergic diseases by differential display of gene showing different expression particularly increased expression in remission in eosinophils of patients, also applicable in screening candidate compounds for remedies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENOX SOYAKU KENKYUSHO KK.
KOKURITSU SHOŅI BYOIN INCHO.
                                                                                                                                                                                         Example 1; Page 56; 72pp; Japanese
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100.001
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11-OCT-2001; 2001WO-JP008937
                                                                                    13-OCT-2000; 2000JP-00314093
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        Unidentified
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method of examining allergic diseases which comprises comparing the expression level of gene B1153 in allergy patients with the expression level in healthy subjects. The method is useful for the treatment, prevention, disagnosis and study of allergic diseases including atopic skin inflammation and asthma. The present sequence is a PCR primer described in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Allergic disease examination method related anchor primer SEQ ID NO: 2.
                                                                                                                                                 Human; allergy; B1153; differential expression; antiallergic; asthma;
antiasthmatic; antiinflammatory; atopic skin inflammation; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                          Examination of allergic diseases comprises detecting gene B1153 over expressed in T cells of allergy patients for diagnosis treatment and investigation of atopic skin inflammation and asthma.
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                                                                                                                              Human B1153 expression in allergic disease related PCR primer GT15A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Allergic disease; allergy; antiallergic; intersectin 2; eosinophil; atopic dermatitis; human; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                        Tsujimoto
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                                                                                                                                                                                                                                                                                                                        Nagasu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                       Sugita Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                           (GENO-) GENOX RES INC.
(NIGE-) JAPAN GEN NAT CHILDREN'S HOSPITAL
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                                                                                                                                                                                                                                                                                                                       Oshida T,
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 2708 TAAAAAAAAAAAAA 2723
                                                                   AAL49948 standard; DNA; 17 BP.
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                                                                                                            (first entry)
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            Best Local Similarity 100.
Matches 16; Conservative
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                                                                                                                                                                                                             WO200250269-A1
                                                                                                                                                                                         Unidentified
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                                                                                                            10-DEC-2002
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                                                                                                                                                                                                                                   Method for examining allergic diseases by differential display of intersectin 2 gene showing different expression particularly significant increase in eosinophils in patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Atopic dermatitis, ss, differential display, primer, PCR; eosinophil; allergic disease; antiallergic; dermatological; GT15a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                             Saito H;
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                                                                                                             Obayashi M, Nagasu T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human atopic dermatitis cDNA related PCR primer GT15a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
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Pred. No. 1.1e+03;
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(NIGE-) JAPAN GEN NAT CHILDREN'S HOSPITAL.
(GENO-) GENOX RES INC.
(NIGE-) JAPAN GEN NAT CHILDREN'S HOSPITAL.
(EISA ) EISAI CO LTD.
                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 52; 90pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; P.
                                                                                                             Ogawa K,
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                                                                                                             Hashida R,
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nes 16; Conserv
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                                                                                                             Sugita Y,
Takahashi E
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This invention relates to gene sequences that are differentially expressed in eosinophils from patients with atopic dermatitis in the increment stage as compared with those in the remission stage. These sequences are used in a novel method for examining allergic diseases comprising determining the expression levels of these genes and comparing individual. The method of the invention may have antiallergic or dermatological activities. The method can be used to diagnose allergic diseases particularly atopic dermatitis, and may also be used to screen candidate compounds for remedies. The method of the invention can be performed in high throughput, at low cost. The present sequence represents the GT15a PCR primer used to amplify the differentially
                                                                                     Examining allergic diseases by differential display of genes showing different expression particularly increase in remission stage in eosinophils in patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amplified atopic dermatitis related cDNA sequences of the invention
    Saito H;
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    Nagasu
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  Fujishima T,
                                                                                                                                                                               Example 1; Page 54; 74pp; Japanese.
  Ogawa K,
Hashida R,
                                           WPI; 2002-330097/36.
Sugita Y,
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ö 0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Indels 2708 TAAAAAAAAAAAA 2723 17 TAAAAAAAAAAAA 16; Conservative Local Similarity Query Match Matches ð g

ADB04271 standard; DNA; 17 BP 20-NOV-2003 (first entry) ADB04271; RESULT 1469 ADB04271 

Human MDZ7 scanning oligonucleotide SEQ ID 5257.

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ1; MDZ12; chromosome 7g22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15g26.1; cancer; developmental disorder; ss.

Homo sapiens

EP1281758-A2

05-FEB-2003

30-JUL-2002; 2002EP-00016874

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC.

Gu Y, Nguyen C; Shannon M,

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 5257; 103pp; English.

proteins and their coding sequences: MD23, MD24, MD21, MD212. MD23 is proteins and their coding sequences: MD23, MD24, MD21, MD212. MD23 is encoded at chromosome 7q22.1, MD24 is encoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15q26.1. The MD24, MD24, MD21 sequences are useful in therapy. Or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring agene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention. The present invention relates to novel human zinc finger-containing

Sequence 17 BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;

ö 0.6%; Score 16; DB 1; Length 17; 0; Indels 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Query Match
Best Local Similarity 100.
Matches 16; Conservative ò

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Gaps

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RESULT 1470

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Gaps

Human MDZ7 scanning oligonucleotide SEQ ID 5258. ADB04272 standard; DNA; 17 BP (first entry) 20-NOV-2003 ADB04272; ADB04272

Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss. 

Homo sapiens

EP1281758-A2

05-FEB-2003.

30-JUL-2002; 2002EP-00016874.

02-AUG-2001; 2001US-00922181

(AEOM-) AEOMICA INC

Shannon M, Gu Y, Nguyen C;

WPI; 2003-423107/40.

New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.

Example 8; SEQ ID NO 5258; 103pp; English.

The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12, MDZ3 is encoded at chromosome 7022.1, MDZ4 is encoded at chromosome 1591.3-22.2, MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing alsorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic

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acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Murine oligonucleotide associated with tumour supression, SEQ ID 2513
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
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                                                                                                                                                                                              DB 1; Length 17; 1.1e+03;
                                                                                                                                                                                                                                  0; Indels
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                                                                                                                                                                                                  0.6%; Score 16;
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                                                                                                                                                                                                                                                                            2709 AAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                               ACC65266 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               schizophrenia; ss.
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New isolated nucleic acid encoding oxalate decarboxylase from Aspergillus phoenices, for degrading oxalic acid, identifying transformed plant cells, and preventing pathogenic disease in plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Aspergillus phoenices; oxalate decarboxylase; APOXD; transgenic plant; crop protection; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0; Indels
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31
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GATCTCTTGGCTGTGG
                                                      GATCTCTTGGCTGTGG
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Best Local Similarity 100.
Matches 16; Conservative
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968
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                                                                                                                                              RESULT
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Gaps

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Query Match 0.6%; Score 16; DB 1; Length 17; Best Local Similarity 100.0%; Pred. No. 1.1e+03; Matches 16; Conservative 0; Mismatches 0; Indels

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Nucleotide sequence of a double stranded product DNA.
                                                                                                                                                                                                                                                       Sequence 17 BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                           represents a double stranded product DNA fragment
                                                                                                                                                                                                                             site usage or determining polyA site usage in a
                                                                                                                                                                                                                                                                                                 2709 AAAAAAAAAAAAA 2724
                                                                                                                    Example, Fig 3, 81pp, English.
                                                                                                                                                                                                                                                                                                                                                ACF36370 standard; DNA; 17 BP
                    28-JAN-2003; 2003WO-IB000255.
                                 29-JAN-2002; 2002US-0352245P
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                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAA 1
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Best Local Similarity 100.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                               (GLOB-) GLOBAL GENOMICS
                                                             Oldin M,
                                                                          WPI; 2003-627619/59
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                                                             Lonnerberg P,
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       07-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                              16
                                                                                                                                                                                                                                                                                                                                                              ACF36370;
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Metsis A, Pihlak A;

Ernfors P, Bauren G,

Linnarsson S,

29-JAN-2002; 2002US-0352215P

(GLOB-) GLOBAL GENOMICS AB

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New naturally derived DNA specifically expressed during blastogenesis of a plant, useful for producing a transformed plant and for compulsive expression of a protein.
                                                                                  Producing a population of double-stranded product DNA molecules, useful for mRNA profiling, comprises amplification by nested polymerase chain reaction.
                                                                                                                                                                                                                         The invention relates to producing a population of double-stranded product DNA molecules comprising amplification by a nested PCR method. The method is useful in profiling mRNA transcribed in a system under investigation. The oligonucleotides are used as size standards in electrophoresis, and as internal controls allowing for calculation of relative amounts of material present. The present sequence represents double stranded product DNA, which aids in outlining an approach to production of a single pattern characteristic of a sample, employing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    during plant blastogenesis. The DNA of the invention is useful for producing a transformed plant. Methods of the invention are also useful for compulsive expression of this DNA. Methods of the invention are useful for plant tissue specific expression of genes. Also, the growth stage of a plant can be controlled specifically. The current sequence represents a PCR primer for amplifying a plant blastogenesis specific gene of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer for amplifying plant blastogenesis specific gene #SEQ ID 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plant blastogenesis; transformation; gene expression; tissue specific;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16; DB 1; Louis Pred. No. 1.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.6%; Scor.
100.0%; Pred. No. 1...
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                                                                                                                                                                                                                                                                                                                                                                                                                      type II restriction enzyme (FokI)
                                                                                                                                                                                     Example; Fig 2; 105pp; English
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                                             WPI; 2003-618365/58.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR; primer; ss
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  Montelius A;
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                                                                                                                                                                                                                                                                                          Determining polyadenylation sites within transcribed gene sequences present in a sample comprises assigning to gene fragments gene candidates within a database by comparing signals in the dataset with the database.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       polyadenylation site within a sequence of a transcribed gene or variants present in a sample. The method involves assigning to gene fragments gene database, the database by comparing signals in the dataset with the database, the database comprising data representing mRNAs with known polyA sites and/or 'virtual genes' representing a possible polyadenylation site within an actual gene. The method is useful for determining the presence of and/or identifying a polyadenylation site or alternative polyadenylation sites within a sequence of a transcribed gene or sequences of transcribed gene fremences of transcribed gene fremences of transcribed gene features, particularly in identifying differences between sequence variants that occur in a population of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             usage or determining polyA site usage in a nucleic acid sample, and variants artising from alternative polyA sites. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to determining the presence of and/or identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid molecules, especially in identifying or discovering polyh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid manipulation; mRNA profiling; polymerase chain reaction; electrophoresis; type II restriction enzyme; FokI; ds.
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Gaps

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The invention relates to a novel isolated nucleic acid that encodes a protein with a chromatin organisation modifier (CHROMO) domain. The polynucleotide of the invention demonstrates cytostatic activity and may be useful for preparing a composition for treating or preventing a composition for treating or preventing a pcoper associated with decreased or increased expression or activity of prost (prostate cancer candidate protein 1), such as a tumour, as well as during gene therapy and vaccine production procedures. The current sequence is that of the human PCCP1-related DNA fragment SEO ID 4-sequence probe of the invention. Note: The current sequence is not shown within the specification per se but was retrieved from the WipoWeb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New prostate cancer candidate protein 1 (PCCP1), useful for preparing a composition for treating or preventing a disorder associated with decreased or increased expression or activity of PCCP1 e.g., tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromatin organisation modifier; CHROMO domain; cytostatic; PCCP1; prostate cancer candidate protein 1; tumour; gene therapy; vaccine;
                                                                                                                                      chromatin organisation modifier; CHROMO domain; cytostatic; PCCP1; prostate cancer candidate protein 1; tumour; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PCCP1 DNA fragment SEQ ID 4-directed probe - SEQ ID 162.
                                                                                               Human PCCP1 DNA fragment SEQ ID 4-directed probe - SEQ ID 161.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.6%; Score 16; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 161; 164pp; English.
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                                                                                                                                                                                                                                                                                                                                                      22-NOV-2002; 2002WO-US037506.
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                                                            12-FEB-2004 (first entry)
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Best Local Similarity 100.
Matches 16, Conservative
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                                                                                                                                                                                          human; ss; probe
                                                                                                                                                                                                                                                                        WO2003050284-A1.
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                                                                                                                                                                                                                                 Homo sapiens
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                   ADF62257;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying novel and useful genes by searching a database utilizing size information about known nucleotide sequences to a specific sequence in a target fragment and the information about the specific sequence to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to novel gene prediction methods. The novel methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprise searching a gene datebase and utilising the information about the size of a known nucleotide sequence to a specific sequence in a target fragment and using the information about the specific sequence in a extract a predicted gene. The methods are used for identifying novel useful genes. The novel gene prediction methods make it possible to predict a gene contained in a DNA fragment obtained as a result of gene expression analysis effectively in a simple and easy manner. The methods and efficiency of genetic analysis is markedly improved. This
                                                                                    Gaps
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100.0%; Pred. No. 1.1e+03;
                                       DB 1; Length 17;
1.1e+03;
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Seguence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
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                                       Query Match 0.6%; Score 16; DB Best Local Similarity 100.0%; Pred. No. 1.1 Matches 16; Conservative 0; Mismatches
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extract a predicted gene.
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nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                               gene prediction;
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18-SEP-2003

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Query Match

Best Local Matches

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RESULT 1477

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28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                The invention relates to a novel isolated nucleic acid that encodes a protein with a chromatin organisation modifier (CHROMO) domain. The polynucleotide of the invention demonstrates cytostatic activity and may be useful for preparing a composition for treating or preventing a disorder associated with decreased or increased expression or activity of PCCP1 (prostate cancer candidate protein 1), such as a tumour, as well as during gene therapy and vaccine production procedures. The current sequence is that of the human PCCP1-related DNA fragment SEQ ID 4-directed probe of the invention. Note: The current sequence is not shown within the specification per se but was retrieved from the WipoWeb
                                                                                                                                                           New prostate cancer candidate protein 1 (PCCP1), useful for preparing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; centerin kinase PKB; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple solerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                       composition for treating or preventing a disorder associated with decreased or increased expression or activity of PCCP1 e.g., tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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Pred. No. 1.1e+03;
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                                                                                    (AMSH ) AMERSHAM BIOSCIENCES SV CORP.
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100.0%; Pro
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                                  22-NOV-2002; 2002WO-US037506.
                                                           10-DEC-2001; 2001US-0339764P
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29-MAY-2001; 2001US-0294412P.
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Best Local Similarity 100.
Matches 16, Conservative
                                                                                                                                   WPI; 2003-532916/50.
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           19-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                             database.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADL48488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1479
                                                                                                            Guo J;
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin DZ receptor (PTGRR),
IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (ONS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), inflammatory disease, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
                                                                                                                                                                                                                       Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autorimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
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                                                                             Fosnaugh
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Pred. No. 1.1e+03;
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                                                                         Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 4 A; 4 C; 7 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                      Claim 59; SEQ ID NO 2021; 317pp; English.
                                                                         Д,
                                                                         Haeberli
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29-MAY-2001; 2001US-0294412P.
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(RIBO-) RIBOZYME PHARM INC.
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nes 16; Conservative
                                                                         Chowrira B,
                                                                                                                                              WPI; 2003-058513/05
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                                                                         3latt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2535
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                                                                                                                                              The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a newrite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
[StappaB kinase [KKK], or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
conditions (e.g. asthma, allergic rhinitis or atopic dermatiis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine generic
diffts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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                                                                                    Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; PCR; ss; allergic disease; NOR-1; MINOR; eosinophil;
atopic dermatitis; antiallergic; antiinflammatory; dermatological;
primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer GT15A used to amplify human NOR-1 (MINOR) DNA SegID 3.
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                                           χ.
                                           Fosnaugh
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1.1e+03;
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                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 5 A; 3 C; 7 G; 0 T; 2 U; 0 Other;
                                           Mcswiggen J,
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                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                               59; SEQ ID NO 2174; 317pp; English
                                                                                                                                                                                                                                                                                                                                                                 0.6%; Score 16;
100.0%; Pred. No.
                                           Haeberli P,
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JAPAN GEN AGENCY NATION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI13009 standard; DNA; 17 BP
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28-AUG-2001; 2001US-0315315P
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                     (RIBO-) RIBOZYME PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 100.
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                                           Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-083057/08.
                                                                 WPI; 2003-058513/05
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                                           Blatt L,
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                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                     Matches
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This invention relates to a novel method for examining allergic diseases that comprises comparing the expression levels of a gene encoding the NOR-1 receptor protein between patients and healthy individuals.

Specifically, the NOR-1 gene, also referred to as MINOR, is expressed in the specialist white blood cells known as eosinophils and is involved in mediating an allergic reaction. The present invention describes a differential display method that can identify the expression level of this gene in order to identify its usefulness in diagnosing allergic diseases such as atopic dermatitis. Furthermore, compositions can also be used to screen compounds for the treatment of allergic diseases. Accordingly, they exhibit various activities including antiallergic.

PER primer used to amplify human NOR-1 DNA in an exemplification of the The invention relates to an oligonucleotide having the formula: (a) 5' XYNIYZN2 3' where 5' denotes the 5' end of the oligonucleotide and 3' denotes the 3' end of the oligonucleotide, where X is a T or modified T nucleotide, X is a G or modified C nucleotide, Z is a G or modified C New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus. Examining allergic diseases e.g. atopic dermatitis by differential display based on gene expression of NOR-1 receptor protein, also applicable in screening compounds for treatment of allergic diseases Gaps pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothloate; ss. . 0 IL-10 expression assay, test oligonucleotide SEQ ID No:33. Length 17; Indels Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other; 0.6%; Score 16; DB 1; Le 100.0%; Pred. No. 1.1e+03; iive 0; Mismatches 0; SEQ ID NO 3; 155pp; Japanese. Example; SEQ ID NO 33; 111pp; English. 2708 TAAAAAAAAAAAAA 2723 (COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH. 묩 04-APR-2005; 2005WO-US011827. 02-APR-2004; 2004US-0558951P. AED81275 standard; DNA; 17 (first entry) 16; Conservative Krieg AM, Vollmer J; WPI; 2005-786756/80. Best Local Similarity WO2005111057-A2 26-JAN-2006 24-NOV-2005. Example 1; Synthetic 11 AED81275; Query Match Matches

connucleotide, and NI and NZ are polymucleotides that do not include a CG dinucleotide, where NI does not include 5' Z nucleotide, and where a 3' polymucleotide consisting of the YZ dinucleotide and the NZ polymucleotide contains a number of nucleotide that is at most 45% of the number of nucleotides in the oligonucleotide, and (b) 5' XYNIYZNZ 3' where 5' denotes the 5' end of the oligonucleotide, where XI as a Tor modified and 3' denotes the 3' end of the oligonucleotide, where XI as a Tor modified T nucleotide, Y is a C or modified C nucleotide, where NI does not include a CG a polymucleotide of 5-10 nucleotides, where NI does not include a CG a polymucleotide of 5 to 30 nucleotides. Also described are: (1) a pharmaceutical composition comprising the oligonucleotide in combination which a therapeutic agent selected from chemotherapeutic agents, monoclonal antibodies, and anticancer agents; (2) a method of specifically increasing interleukin (IL)-10 expression and antiche comprising contraction of the services and anticancer agents. distriction to interior the control time a subject, comprising an oligonaclectide or a pharmaceutical composition to the subject in need of increased IL-10 expression relative to IFN-alpha expression, (3) a method of inducing an antigen-specific regulatory T call response in a subject by administering an immunostimulatory nucleic exposed to an antigen, (4) a method of inducing an antigen-specific regulatory B cell response in a subject by administering an antigen-specific regulatory B cell response in a subject by administering an antigen, (5) a method of treating an allergy or subject exposed to an antigen, (5) a method of treating an allergy or subject exposed to an antigen, (5) a method of treating an allergy or immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an anount sufficient to prevent or alleviate an allergic response to the allergen in the subject; (6) a method of treating an autoimmune disease to an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an immunostimulatory nucleic acid or composition is administered in an acount sufficient to prevent or treat an autoimmune disease in the subject; and (7) a method of reducing an antigen-specific response to an implant in a subject by exposing a subject to an implant antigen, and anti subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigenspecific response to the implant in the subject. The oligonucleotide includes at least 1 modified internucleotide linkage such as a phosphorothioate linkage. The oligonucleotide, methods and compositions of the invention are useful for treating allergies, asthma, autoimmune diseases, arthritis, systemic lupus erythemacosus, multiple sclerosis, crohn's disease, Type 1 diabetes mellitus, myasthenia gravis, autoimmune neuropathies such as Guillain-Barre, autoimmune uveltis, autoimmune hemolytic anemia, pernicious anemia, autoimmune thrombocytopenia, Crohn's polymyositis, dermatcomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, vitiligo, Crohn's Disease, ulcerative colitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated diabetes mellitus, Grave's Disease, Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma, invention.

Sequence 17 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 1 Other;

; 0 0.6%; Score 16; DB 1; Length 17; 14.1%; Pred. No. 1.1e+03; ve 0; Mismatches 1; Indels Query Match 0.6%; Best Local Similarity 94.1%; Matches 16; Conservative

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Gaps

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AED81272/

AED81272 standard; DNA; 17 BP. RESULT 1483

AED81272;

(first entry) 26-JAN-2006

IL-10 expression assay, test oligonucleotide SEQ ID No:30.

pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.

Synthetic.

WO2005111057-A2.

24-NOV-2005.

04-APR-2005; 2005WO-US011827.

)2-APR-2004; 2004US-0558951P.

(COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH.

Vollmer J; Krieg AM, WPI; 2005-786756/80.

New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type I diabetes mellitus.

Example, SEQ ID NO 30; 111pp; English.

The invention relates to an oligonucleotide having the formula: (a) 5, XYNIXZN2 3, where 5, denotes the 5, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide, where X is a T or modified T uncleotide, Y is a C or modified C nucleotide, and or modified G uncleotide, and where a 3, or uncleotide, where N is a C or modified C nucleotide and the N2 mucleotide, and where a 3, polynucleotide, where N is number of nucleotide and the N2 comparing of the YZ dinucleotide, and where a 3, polynucleotide consisting of the YZ dinucleotide and the N2 polynucleotide consisting of the PZ dinucleotide and the N2 polynucleotide consisting of the Oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of the oligonucleotide and 3, denotes the 3, end of 5 to 3 nucleotides, where N1 does not include 5, Z nucleotide, N1 is a polynucleotide of 5 to 3 nucleotides. Also described are: (1) a polynucleotide of 5 to 3 nucleotides. Also described are: (1) a paramaceutical composition comprising the oligonucleotide in combination with a therapeutic agents, monoclonal antibodies, and anticancer agents; (2) a method of specifically increasing interleuxin (11)-10 expression relative to interferen (17)-10 expression relative to interferen (17)-10 expression relative to interferen (17)-10 expression relative to onterferen (17)-10 expression relative to or interferent (17)-10 expression relative to interferent (17)-10 expression relativ amount sufficient to prevent or alleviate an allergic response to the allergen in the subject; (6) a method of treating an autoimmune disease in a subject by exposing a subject to a self antigen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or treat an autoimmune disease in the subject; and (7) a method of reducing an antigen-specific response to an implant in a subject by exposing a subject to an implant antigen, and administering an immunostimulatory nucleic acid or composition to the

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administered in an amount sufficient to prevent or reduce an antigen-
specific response to the implant in the subject. The oligonucleotide
includes at least I modified internucleotide linkage such as a
phosphorothicate linkage. The oligonucleotide, methods and compositions
of the invention are useful for treating allergies, asthma, autoimmune
diseases, arthritis, systemmic lupus erythemacosus, multiple sclerosis,
crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune
hemolytic anemia, pernicious anemia, autoimmune thromborytopenia,
cemporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus
vulgaris, vasculitis such as Wegener's granulomatosis, vitiligo, Crohn's
Disease, ulcerative colitis, primary biliary cirrhosis, autoimmune
hepatitis, immune-mediated diabetes mellitus, Grave's Disease,
Hashimoto's thyroiditis, autoimmune opphoritis and orchitis, autoimmune
clisease of the adrenal gland, rheumatoid arthritis, scleroderma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA fragment coding for Papillomavirus antigenic proteins - and derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
the immunostimulatory nucleic acid or composition is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ll region of the bovine papillomavirus type la genome, fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diagnostioc reagent; vaccine; medicine; wart; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 1 Other;
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1. .18
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunogen, vaccine and antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2709 AAAAAAAAAAAAAA 2725
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Best Local Similarity 94.1%;
Matches 16; Conservative
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Unidentified.
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P-PSDB; AAP30313.
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05-APR-1992
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      The inventors claim DNA fragments capable of expressing, in a host, a prod. contg. at least one antigenic determinant of papillomavirus (PV), (See AAN30170-01731). Also claimed are immunogens consisting of at least one peptide sequence coded for by the DNA fragments (see AAP30310-P30313), vaccines contg. the immunogens and antibodies raised from them. antibodies are useful in human and veterinary medicine and the antibodies are useful as diagnostic reagents. The DNA fragments are most esp. derived from the L1 region of human PV type la
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                                                                                                                                                Revised record issued on 09-SEP-2004 : Correction of feature table key
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                                                                                                                                                                             Sequence 18 BP; 16 A; 1 C; 1 G; 0 T; 0 U; 0 Other;
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100.0%; Pred. No. 1.1e+03;
ive 0; Mismatches 0;
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Pred. No. 1.1e+03;
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100.001
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Matches
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Synthetic.

AAV54164;

AAV54164

Sakaki Y;

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This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperuricemia and sleep apnea syndrome. The genes (AA296631-633) and the proteins (AAV67598-V67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AA29640-51 represent PCR primers amplifying the human adipose tissue genes
                                                                                                                                                                                Novel apoptosis-related DNAs and proteins - for diagnosis, preventing or treating diseases associated with apoptosis.
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Pred. No. 1.1e+03;
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AAZ90649 standard; DNA; 18
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                                                                                                          Sakaki Y;
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Matches 16; Conservative
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Sequence 18 BP; 1 A; 1 C; 1 G; 15 T; 0 U; 0 Other;

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Query Match

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Homo sapiens.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                   A physiologically active protein specifically derived from mammal tissue
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100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0; Indels
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1.1e+03;
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                                                                                                                                                                           Human adipose tissue gene amplifying primer #7.
        0.6%; Score 16; DB
100.0%; Pred. No. 1.1
ive 0; Mismatches
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                                                                                                                  AAZ90646 standard; DNA; 18 BP
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    Query Match
Best Local Similarity 100.0
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                                                2708 TAAAAAAAAAAAAA
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Best Local Similarity 100.
Matches 16; Conservative
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                                                                                                                                                                                                                             Homo sapiens
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A physiologically active protein specifically derived from mammal tissue.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Pred. No. 1.1e+03;
0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 100; 101pp; English
                                                                                                                                                                                                                                                                                                           Example 2; Page 18; 50pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Pre-
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06-APR-2000; 2000US-00544713.
98JP-00225228
                                                             98JP-00225228
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                                                                                                                       (NISB ) JAPAN TOBACCO INC.
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nes 16, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-202878/20.
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Gaps

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Lags from nucleic acid fragments in a sample, involving binary sequence tags from nucleic acid fragments in a sample, involving incubating the sample with cleaving reagents, mixing offset adaptors with the sample, incubating with more cleaving reagents and mixing the sample with adaptor -indexers where the adaptors are coupled to binary sequence tags. The method is useful in sequence analysis, including analysis and comparison of gene expression, nucleic acid samples and genomes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for treating and/or preventing cytomegalovirus (CMV) infection and/or related diseases. The method of the invention comprises contacting a test compound with at least one of the cellular kinases RICK, RIP, Nck-Interacting kinase, MKK3 and SPRY-2 and detecting any change in kinase activity. The method of the invention can be used to treat and/or prevent CMV infections and related diseases. Oligonucleotides that can detect the specified kinases can also be used for diagnosis of infection. The present nucleic acid sequence represents human CMV reverse transcriptase (RT)-PCR primer TXN that was used in the methods of the invention for preparation of radioactively labelled cDNA probes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention relates to a new method for identifying compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  compound with cellular kinase and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cytomegalovirus; HCMV; virucide; cytomegalovirus infection;
cellular kinase; RICK; RIP; Nck-Interacting kinase; MKK3; SRPK-2;
reverse transcriptase PCR; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytomegalovirus
                                                                                                                                                                                                        0.6%; Score 16; DB 1; Length 18; 100.0%; Pred. No. 1.1e+03; ive 0; Mismatches 0; Indels
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                                                                                                                                                                     Sequence 18 BP; 0 A; 1 C; 1 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human cytomegalovirus (HCMV) RT-PCR primer TXN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection, comprises contacting test compound detecting change in cellular kinase activity.
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                                                                                                                                                                                                                                                                                2709 AAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                ABK51158 standard; DNA; 18 BP.
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                                                                                                                                                                                                                                                                                                                   16 AAAAAAAAAAAA 1
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                                                                                                                                                                                                                                             16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cytomegalovirus
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                                                                                                                                                                                                          Query Match
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The invention relates to novel pyridylpyrimidine derivatives and methods of detecting prion infections and/or prion disease in an individual or in cells, cell cultures and/or cell lysates. The method involves adding at least one monoclonal or polyclonal antibody, oligonucleotide or pyridyleast one monoclonal or polyclonal antibody, oligonucleotide or pyridylex pyrimidine derivative to the sample or in cells, cell cultures and/or cell lysates and detecting the activity of at least one human cellular protein kinases (e.g., FGF-R1 (also known as flg, F1-1, F1t-2, b-FGFR). Tkt (also known as CCK-2, DDR-2 or EDDR; EC number 2.7.1.112), Abl (also known as CCK), PRK, human cellular protein phosphatases such as PTP-SL known as CCK1), PRK, human cellular protein phosphatases such as PTP-SL (also known as MCP83) and PTP-zeta, the cellular signal transduction of prions in cells and in the manufacture of pharmaceutical production of prions in cells and in the manufacture of pharmaceutical composition for prophylaxis and/or treatment of infectious disease (e.g. composition for prophylaxis and/or treatment of infectious disease (e.g. CCR production of prions in cells and in the manufacture of pharmaceutical composition wasting disease (CMD), transmissible mink encephalopathy (TMB), creutzfeldt-Jacob disease (CMD), bovine spongiform encephalopathy (EME), variant CUD, Gerstmann-Straussler-Scheinker syndrome (GSS), fatal familial insomnia (FFI), Kuru and Alpers syndrome, especially BSE, CUD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; pyridylpyrimidine derivative; cellular protein kinase; Scrapie; cellular protein prophylaxis; protein phosphatase; cellular signal transduction; prophylaxis; prion infection; chronic wasting disease; CWD; Creuzfeldt-Jacob disease; CJD; transmissible mink encephalopathy; bovine spongiform encephalopathy; TME; SBE; Gerkemann-Straussler-Scheinker syndrome; GSS; Alpers syndrome; fatal familial insomnia; FPI; kuru; neurodegenerative disease; nootropic; Alzheimer's disease; primer; skuru;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New pyridylpyrimidine derivatives useful in the treatment or prevention of infectious disease e.g. Kuru syndrome and Creutzfeld-Jacob disease
                                                                                                            Gaps
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                                                               Score 16; DB 1; Length 18;
Pred. No. 1.1e+03;
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                    Sequence 18 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 2 Other;
                                                        0.6%; ccc
100.0%; Pred. No. ...
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AAD52799 standard; DNA; 18 BP.
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29-MAY-2001; 2001US-0293528P.
13-JUL-2001; 2001EP-00117113.
18-JUL-2001; 2001US-0305898P.
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                                                                                                                                                                           (first entry)
                                                                                                       16; Conservative
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                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200293164-A2.
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                                                            Query Match
                                                                                                                                                                                                                                                         RESULT 1493
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The invention relates to an anti-proliferative substance for preventing uncontrolled cellular proliferation comprising a radiolabelled DNA carrier, where a radiolabelled DNA carrier, where a radiolabelled DNA carrier aguence, at 5' end or at 3' end, and the radiolabelled DNA carrier penetrates the cell membrane and is retained intracellularly for a time in the anti-proliferative substance is an oligonucleotide, which is linear or a plasmid, which is circular. The plasmid is of viral or bacterial origonucleotide is a double- or a single-stranded DNA sequence, which is conjugated with an antibody for cell-specific delivery. The oligonucleotide is also conjugated to a stent surface, choice acid, losic acid, 
                                                                                                                                                                                                                                      ö
vCJD) or neurodegenerative diseases (e.g., Alzheimer's disease) in humans or ruminants. The present DNA sequence is a primer used to prepare radioactively labelled cDNA probes from RNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98; anti-proliferative; cellular proliferation; restenosis; angioplasty;
cancer; malignant tumour.
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                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                    Length 18;
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                                                                                                                                Seguence 18 BP; 0 A; 0 C; 0 G; 16 T; 0 U; 2 Other;
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1.1e+03;
                                                                                                                                                                                 0.6%; Score 16; DB 100.0%; Pred. No. 1.1 tive 0; Mismatches
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99US-00318106
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Best Local Similarity 100.0
Matches 16; Conservative
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24-MAY-1999;
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The invention relates to an antisense oligonucleotide or its active derivative selected from AEC46395, targeting human interleuking derivative selected from AEC46374-AEC46395, targeting human interleuking control of [1L-10]. Also included are a process of manufacturing the antisense coligonucleotide (or its active derivative, by adding consecutive nucleosides and linker stepwise or by cutting the oligonucleotide out of nucleosides and linker stepwise or by cutting the oligonucleotide out of composition for treating cancer. The oligonucleotide is an antisense oligonucleotide inhibiting the synthesis of proteins involved in the composition of metastases. The oligonucleotide is an antisense oligonucleotide inhibiting the production of transforming growth factor (TGF)-beta 1, TGF-beta 3, TGF-beta 3, call-cell adhesion molecules (CMMS), integrins, selectins, metalloproteases (MMPs), their tissue confinibitors (TTMPs) and/or interleukins 10. The oligonucleotides are useful for the preparation of a pharmaceutical composition for inhibiting the formation of metastases in cancer treatment. The oligonucleotides are useful for treating cancer, bronchogenic carcinoma, carcinoma, cervical carcinoma, colon carcinoma, colon carcinoma, coloneactal carcinoma, embryonal carcinoma,
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uncontrolled cell proliferation is a restenosis following angioplasty,
                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides inhibiting the synthesis of proteins involved in the formation of metastases such as transforming growth factor-beta 1 (TGF-beta 1), TGF-beta 2 and TGF-beta 3, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stauder G;
               cancer or a malignant tumour. The present sequence represents an oligonucleotide carrier used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transforming growth factor beta; TGF-beta-3; antisense therapy; antisense oligonucleotide; ss; cancer; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide targeting human TGF-beta-3 #871.
                                                                                                               Length 18;
                                                                                                                                                0; Indels
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                                                                          Sequence 18 BP; 3 A; 0 C; 0 G; 15 T; 0 U; 0 Other;
                                                                                                               DB 1; Le
1.1e+03;
                                                                                                                          100.0%; Prec. ...
                                                                                                               Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schlingensiepen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 71; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schlingensiepen K, Schlingensionschof A, Hafner M, Egger T;
                                                                                                                                                                                         2708 TAAAAAAAAAAAA 2723
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01-APR-2004; 2004US-0558135P.
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                                                                                                                                                                                                                              16 TAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                         AEC52473 standard; DNA; 18
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                                                                                                                                Local Similarity 100.
nes 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                              AEC52473;
                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                      RESULT 1495
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endometrial cancer, epithelial carcinoma, esophageal cancer, gall bladder cancer, logatric cancer, head and neck cancer, hepatocellular cancer, liver carcinoma, lung carcinoma, medulary carcinoma, non-small cell bronchogenic/lung carcinoma, medulary carcinoma, papillary carcinoma, papillary adenocarcinoma, prostate cancer, small intestine carcinoma, rectal cancer, renal cell carcinoma, sebaceous gland carcinoma, skin cancer, soft tissue cancer, squamous cell carcinoma, testicoma, uterine cancer, squamous cell carcinoma, trachomas, and pyogenic granulomas; pre-malignant tumors, blastoma, bwing's tumor, craniopharyngioma, ependymoma, medulloblastoma, neurofibroma, neurofibroma, medullablastoma, medullablastoma, medullablastoma, neurofibroma, retinoblastoma, sarcoma, seminoma, trachomas, neurofibroma, pinealoma, retinoblastoma, sarcoma, seminoma, trachomas, anticone oli cancer, successivant cancer, seminoma, cancer, cancer, cancer, seminoma, cancer, cancer, cancer, cancer, cancer, cancer, cancer, cancer, can
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 16; DB 1; Length 18;
100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0; Indels
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M, Egger T;
                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotide targeting human TGF-beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 15 A; 0 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transforming growth factor beta; TGF-beta-3; antisantisense oligonucleotide; ss; cancer; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ANTI-) ANTISENSE PHARMA GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AEC52193 standard; DNA; 18 BP
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01-APR-2004; 2004US-0558135P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16; Conservative
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Bischof A, Hafner M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-SEP-2005.
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Control of the case and the production of transforming growth factor oligonucleotide inhibiting the production of transforming growth factor (TGF)-beta 1, TGF-beta 2, TGF-beta 3, cell-cell adhesion molecules (CDMS), integrins, selectins, metalloprofeases (MMPB), their tissue inhibitors (TIMPB) and/or interleukins 10. The oligonucleotides are useful for the preparation of a pharmaceutical composition for inhibiting the formation of metastases in cancer treatment. The oligonucleotides are useful for treating cancer, e.g. bile duct carcinoma, bladder carcinoma, brain tumor, breast cancer, bronchogenic carcinoma, carcinoma of the carcinoma, colon carcinoma, colorectal carcinoma, colorectal carcinoma, encounted a carcinoma, colorectal carcinoma, emphysonal carcinoma, endometrial cancer, epithelial carcinoma, esophageal cancer, gall bladder carcinoma, lung carcinoma, medullary carcinoma, non-small cell cancer, gastric cancer, epithelial carcinoma, medullary carcinoma, non-small cell carcinoma, papillary carcinoma, medullary carcinoma, papillary adenocarcinoma, prostate cancer, small coll carcinoma, sebaceous gland carcinoma, skin cancer, soft tissue cancer, squamous carcinoma, terinoma, uterine cancer, soft tissue cancer, squamous carcinoma, tersinoma, uterine cancer, soft tissue cancer, squamous carcinoma, tersinoma, uterine cancer, acustic neuromas, neurofibromas, trachomas, and pyogenic granulomas; pre-malignant tumors, blastoma, Ewing's tumor and/or myeloma, medullablastoma, mesothelioma, neuroblastoma, neurofibroma, timeslens pincaloma, retinoblastoma, mesothelioma, seminoma, trachomas, antisone lingual retareine carcinoma, retinoblastoma, mesothelioma, seminoma, trachomas, antisone pincareine carcinoma, retinoblastoma, mesothelioma, seminoma, trachomas, antisone pincareine carcinoma, medullablatoma, mesothelioma, seminoma, trachomas, antisone pincareine carcinoma, medullablatoma, metallablatoma, mesothelioma, seminoma, trachomas, antisone pincareine carcinoma, metallablatoma, metallablatoma, metallablatoma, mes Gaps New antisense oligonucleotides inhibiting the synthesis of proteins involved in the formation of metastases such as transforming growth factor-beta 1 (TGF-beta 1), TGF-beta 2 and TGF-beta 3, useful for ö Transforming growth factor beta; TGF-beta-3; antisense therapy; antisense oligonucleotide; ss; cancer; cytostatic. ö formation of metastases. The oligonucleotide is an antisense Antisense oligonucleotide targeting human TGF-beta-3 #731. 0.6%; Score 16; DB 1; Length 18; 100.0%; Pred. No. 1.1e+03; 0; Indels Schlingensiepen K, Schlingensiepen R, Jachimczak P, Bischof A, Hafner M, Egger T; antisense oligonucleotide targeting human TGF-beta-3 Sequence 18 BP; 15 A; 0 C; 0 G; 3 T; 0 U; 0 Other; 100.0%; Pred. No. 2708 TAAAAAAAAAAAA 2723 (ANTI-) ANTISENSE PHARMA GMBH. AEC52333 standard; DNA; 18 BP 28-FEB-2005; 2005WO-EP002101. 27-FEB-2004; 2004EP-00004478. 01-APR-2004; 2004US-0558135P. (first entry) 16; Conservative Similarity WPI; 2005-630685/64. WO2005084712-A2. treating cancer Homo sapiens. 17-NOV-2005 15-SEP-2005. AEC5233; Query Match Local RESULT 1497 Matches ð 셤 

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ADR81681
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                       The invention relates to an antisense oligonuclectide or its active

derivative selected from AEC46374-AEC46395, targeting human interleukin-
C derivative selected from AEC46374-AEC46395, targeting human interleukin-
C 10 (11-10). Also included are a process of manufacturing the antisense
oligonuclectide chain, a pharmaceutical composition comprising
the antisense oligonuclectide chain, a pharmaceutical composition comprising
the antisense oligonuclectide and TGF-beta 2 antagonist for preparing
c composition for treating cancer. The oligonuclectide is an antisense
oligonuclectide inhibiting the synthesis of proteins involved in the
formation of metastases. The oligonuclectide is an antisense
composition for treating the production of transforming growth factor
(TGF)-beta 1, TGF-beta 2, TGF-beta 3, cell-cell adhesion molecules
c (TGMS), integrins, selectins, metalloproceases (MMPs), their tissue
inhibitors (TIMPs) and/or interleukins 10. The oligonuclectides are
c (TGMS), integrins, selectins, metalloproceases (MMPs), their tissue
c (TMMS), integrins, selectins, metalloproceases (MMPs), their tissue
c carcinoma, colon carcinoma, colorectal carcinoma, carcinoma of the
carcinoma, colon carcinoma, colorectal carcinoma, embryonal carcinoma,
carcinoma, lung carcinoma, medullary carcinoma, papillary carcinoma, medullary carcinoma, papillary daedencearcinoma, procease acarcinoma,
papillary carcinoma, papillary adencer, renal cell carcinoma, papillary carcinoma, papillary daedencer, senter, sender, senter, senter, senter, senter, carcinoma,
papillary carcinoma, uterine cancer, renal cell cancer, senal
trestine carcinoma, uterine cancer, senders, acument, medullatoma, medullato
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hemangioblastoma, medullablastoma, melanoma, mesothelioma, neuroblastoma, neurofibroma, pinealoma, retinoblastoma, sarcoma, seminoma, trachomas, Wilm's tumor and/or myeloma, multiple. The present sequence is an
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e regulation; gene expression; ss; tag.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5' mRNA DNA preparation method related tag DNA sequence #30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.6%; Score 16; DB 1; Length 18;
100.0%; Pred. No. 1.1e+03;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide targeting human TGF-beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 15 A; 0 C; 0 G; 3 T; 0 U; 0 Other;
 Claim 4; Page 71; 106pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2708 TAAAAAAAAAAAA 2723
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12-AUG-2002; 2002JP-00235294.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene regulation;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; anticonvulsant; notropic; muscula; anti-HIV;
RNA interference; iRNA, antisense technology; lipid metabolism;
coholesterol imbalance; dyslipidaemia hypercholesterolaemia;
coronary artery disease; CAD; coronary heart disease; CHD;
atherosclerosis; hepatic glucose production;
glucose-metabolism-related disorder; disease; cancer; breast cancer;
colon cancer; lung cancer; neurological disease; Huntington disease;
spinocerebellar ataxia; viral disease; ADDS; hepatitis C virus; HCV; ss.
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                                                                                                                                                                                                      Preparing DNA fragment corresponding to nucleotide sequence of 5' end region of mRNA, by preparing nucleic acid corresponding to nucleotide sequence of 5' end of mRNA, cleaving nucleic acid with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 16 A; 2 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       Example 5; SEQ ID NO 62; 121pp; English.
                                                                     Harbers MT;
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13-MAR-2003; 2003US-0454262P.
13-MAR-2003; 2003US-0454962P.
13-MAR-2003; 2003US-0455050P.
14-APR-2003; 2003US-0462894P.
17-APR-2003; 2003US-0465802P.
25-APR-2003; 2003US-0465802P.
09-MAY-2003; 2003US-0469612P.
08-MUG-2003; 2003US-04993986F.
11-AUG-2003; 2003US-0494597P.
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                                                                     Carninci P,
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                                                                                                                                    WPI; 2004-082194/08.
(DNAF-) DNAFORM KK.
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                                                                     Hayashizaki Y,
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                                                                                                                                                                                                                                                                                                                            enzyme.
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The invention describes a RNA interference (iRNA) agent (1) comprising a sense sequence and an antisense sequence, where the sense sequences have one or more asymmetrical 2'-0 alkyl modifications, the antisense captuences have one or more asymmetrical phosphorothicate modifications and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (1); reducing (M1) apoB-100 levels or glucose-6-phosphatase levels in a subject; producing (II); introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (1) and instruction for its use; and a device the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (1) and instruction for its use; and a device the modification decreases nuclease sensitivity while not decreased to useful for reducing apoB-100 levels or glucose-6-phosphatase levels.

Cuseful for reducing apoB-100 levels or glucose-6-phosphatase levels.

The subject is suffering from a disorder characterised by elevated or otherwise unwanted expression of apos-100, elevated or otherwise unwanted clevels or disorder cholesterol, and/or disregulation of lipid metabolism. The disorder is chosen from the HDL/LDL cholesterol imbalance, dyslipidaemias, hypercholesterolaemia, statin-resistant chypercholesterolaemia, coronary attery disease (CHD), coronary heart disease (CHD) and atherosclerosis. (1) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism-crelated diseases as mentioned above, cancer (e.g. breast, colon or treating the diseases as mentioned above, cancer (e.g. breast, colon or submocerabellar ataxia) or viral disease (e.g., AIDS). This sequence represents a hepatitie C virus (HCV) antisense oligonucleotide that cancer, especial disease (e.g., AIDS). This sequence
                                                                                                                                                                                                                                                                         Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense sequence and antisense sequence which has specific modifications.
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  26-SEP-2003; 2003US-0506341P.
09-0CT-2003; 2003US-0510246P.
10-0CT-2003; 2003US-0510318P.
07-NOV-2003; 2003US-0518453P.
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16; Conservative
                                                                                                                                                                                 Bumcrot D;
                                                                                                                             (ALNY-) ALNYLAM PHARM.
                                                                                                                                                                                                                             WPI; 2004-677362/66.
26-SEP-2003;
                                                                                                                                                                                 Manoharan M,
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Matches
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ADT86138 standard; DNA; 19 BP (first entry) 13-JAN-2005 ADT86138; 

Hepatitis C virus (HCV) antisense inhibition target segid 6180.

antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic; interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase; lipid metabolism; cholesterol imbalance; dyslipidaemia; familial combined hyperlipidaemia; acquired hyperlipidaemia; hypercholesterolaemia; statin-resistant hypercholesterolaemia; coronary attery disease; coronary heart disease; atherosclerosis;

hepatic glucose production, glucose-metabolism-related disorder, type-2 diabetes, glitaxzone-resistant diabetes, HCV, hepatitis C virus, 10-APR-2003; 2003US-0461915P.
11-APR-2003; 2003US-0461915P.
11-APR-2003; 2003US-046194P.
25-APR-2003; 2003US-0465802P.
25-APR-2003; 2003US-0465802P.
09-MAY-2003; 2003US-0469612P.
09-MAY-2003; 2003US-0499986P.
11-AUG-2003; 2003US-0499597P.
26-SEP-2003; 2003US-0499597P.
26-SEP-2003; 2003US-0494597P.
10-OCT-2003; 2003US-0510318P.
07-NOV-2003; 2003US-0510318P.
08-MAR-2004; 2004WO-US0037070. 09-APR-2004; 2004WO-US011255 05-APR-2004; 2004WO-US010586 type-2 diabetes; glitaxzo antisense inhibition; ss. Hepatitis C virus. WO2004091515-A2 28-OCT-2004. 

(ALNY-) ALNYLAM PHARM INC

Harborth J; Elbashir S, Manoharan M,

WPI; 2004-766693/75.

New interference RNA agent comprising sense sequence and antisense sequence having cholesterol moieties, useful for reducing apoB-100 levels glucose-6-phosphatase levels.

Example 4; SEQ ID NO 6180; 324pp; English.

Computes one of more contractory materies, and the contractor of the contractory of the contractory of the contractory of the contractor o artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesteroledemia. (I) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism-related disorders e.g., type-2 diabetes or glitaxzone-resistant diabetes. (I) has endonuclease or exonuclease resistance. This sequence represents a human hepatitis C virus (HCV) pallindromic sequence that may be useful as a target for antisense inhibition of HCV in human liver cells. The invention describes an interference RNA (iRNA) agent (I) comprising sense sequence and an antisense sequence, where the sense sequence comprises one or more cholesterol moieties, and the antisense sequence

Sequence 19 BP; 0 A; 0 C; 2 G; 17 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 16; DB 1; Length 19; 100.0%; Pred. No. 1.2e+03; tive 0; Mismatches 0; Indels Local Similarity 100. es 16; Conservative Query Match Matches

2709 AAAAAAAAAAAAA 2724

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19 AAAAAAAAAAAAAA

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The invention relates to a method (MI) for treating spinal cord injury in a subject. (MI) involves administering to the subject, a short interfering nucleic acid (siNA) molecule (I) that directs cleavage of a Pas RNA through RNA interference (RNAi) under conditions suitable to modulate the expression of Fas in the subject. Also described: (I) an expression vector comprising (I); (2) a kit comprising (I); (3) a human cell comportaing (I); (4) a pharmaceutical composition comprising (I); and fas gene and Fas ligand (Pasl) lower (antisense) siRNA oligonnoleotide, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating spinal cord injury in subject, involves administering to subject, short interfering nucleic acid directing cleavage of Fas RNA through RNA interference under conditions suitable to modulate expression
                                                                                                                                                                                                                                                                            spinal cord injury; short interfering RNA; siRNA; RNA interference; gene silencing; RNA cleavage; Fas; vulnerary; ds.
                                                                                                                                                                                                                        Human Fas and FasL genes lower siRNA sequence SEQ ID NO:300.
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2003US-00693059.
2003US-00720448.
2004US-0075780.
2004US-0075780.
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2003WO-US005028.
2003WO-US005346.
2003US-00427160.
                                                      AEA99200 standard; RNA; 19 BP
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2002US-0363124P
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                                                                                                                                                                 11-AUG-2005 (first entry)
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29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
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10-FEB-2004;
13-FEB-2004;
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20-MAY-2002;
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RESULT 1501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a method (M1) for treating spinal cord injury in a subject. (M1) involves administering to the subject, a short interfering nucleic acid (siNA) molecule (1) that directs cleavage of a
                                                           Gaps
                                                                                                                                                                                                                                                                       spinal cord injury; short interfering RNA; siRNA; RNA interference; gene silencing; RNA cleavage; Fas; vulnerary; ds.
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                                                                                                                                                                                                                                                 Human Fas and FasL genes target and upper siRNA SEQ ID NO:150
                                   Length 19;
                                                          Indels
         Sequence 19 BP; 2 A; 2 C; 0 G; 0 T; 15 U; 0 Other;
                                0.6%; Score 16; DB 1; Ld
100.0%; Pred. No. 1.2e+03;
tive 0; Mismatches 0;
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2002US-0363124P.
2002WO-US015876.
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2003US-00693059.
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03-DEC-2003; 2003US-00727780.
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2004WO-US013456
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                                                                                                                                                                                                                          (first entry)
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                                              Local Similarity 100.
ses 16; Conservative
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15-JAN-2003;
20-FEB-2003;
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30-APR-2003;
23-MAY-2003;
23-OCT-2003;
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11-MAR-2002;
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                                    Query Match
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Fas RNA through RNA interference (RNAi) under conditions suitable to modulate the expression of Fas in the subject. Also described: (1) an expression vector comprising (1); (2) a kit comprising (1); (3) a human cell comprising (1); (4) a pharmaceutical composition comprising (1); and (5) a method of synthesizing (1). The present sequence represents a human Fas gene and Fas ligand (FasL) target and upper (sense) siRNA oligoniclectide, which is used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        intercellular adhesion molecule 1, ICAM1, CNS-Gen.; Neuroprotective; Notropic; antinflammatory; Antiarthritic; Antirheumatic; Antiasthmatic; Gastrointestinal-Gen.; Dermatological; Immunosuppressive; Antidiabetic; Cytostatic; Cerebroprotective; Respiratory-Gen.; Hypotensive; inflammation, rheumatoid arthritis; inflammatory bowel disease; atopic dermatitis; asthma; autoimmune disease; multiple sclerosis; crohns disease; diabetes mellitus; cancer; hyperproliferation; neurological disease; Alzheimers disease; brain injury; myopathy; respiratory disease; chronic obstructive pulmonary disease; pulmonary hypertension; emphysema; muscular-gen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of intracellular adhesion molecule RNA through RNA interference, useful for treating diseases e.g. cancer and
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                                                                                                                                                                                            0.6%; Score 16; DB 1; Length 19;
13.8%; Pred. No. 1.2e+03;
ve 1; Mismatches 0; Indels
                                                                                                                                                            Sequence 19 BP; 15 A; 0 C; 2 G; 0 T; 2 U; 0 Other;
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2004US-00757803.
2004US-0543480P.
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2004US-00800487.
2004US-00826966.
2004WO-US013456.
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                                                                                                                                                                                                                                15; Conservative
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                                                                                                                                                                                                              Local Similarity
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Claim 33; SEQ ID NO 328; 200pp; English.

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The invention relates to a chemically synthesized double-stranded short interfering nucleic acid (sinA) molecule directing cleavage of interfering nucleic acid (sinA) molecule directing cleavage of interference (RNA1), where each strand of the molecule is 18-23 nucleotides in length, and one strand of the molecule comprises nucleotide sequence having sufficient complementarity to the ICAM RNA for the sinA molecule to direct cleavage of the ICAM RNA through RNA. The sinA is useful for downregulation or inhibition of expression of ICAM gene and for diagnosing, preventing or treating diseases associated with cellular adhesion such as inflammatory disorders e.g. rheumatoid arthritis, inflammatory bowel disease, atopic dermatitis and asthma, autoimmune cancer and proliferative diseases e.g. ovarian cancer, lung cancer, renal cancer and proliferative diseases e.g. ovarian cancer, lung cancer, renal cancer and proliferative diseases e.g. ovarian cancer, lung cancer, renal cancer and sisease, brain injury, cerebral atrophy and congenital myopathy, respiratory diseases e.g. chronic obstructive pulmonary hypertension and emphysema (many more diseases are given in the specification). The present sequence one strand of an siRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    intercellular adhesion molecule 1; ICAM1; CNS-Gen.; Neuroprotective; Notropic; antiinflammatory; Antiarthritic; Antirheumatic; Antiasthmatic; Gastrointestinal-Gen.; Dermatological; Immunosuppressive; Antidiabetic; Cytostatic; Cerebroprotective; Respiratory-Gen.; Hypotensive; inflammation; rheumatoid arthritis; inflammatory bowel disease; atopic dermatitis; asthma; autoimmune disease; multiple sclerosis; Crohns disease, diabetes mellitus; cancer; hyperproliferation; neurological disease; Alzheimers disease; brain injury; myopathy; respiratory disease; chronic obstructive pulmonary disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA interference; gene silencing; siRNA; short interfering RNA; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 16 A; 1 C; 1 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16; DB 1; Le
Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pulmonary hypertension; emphysema; muscular-gen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human ICAM1 siRNA lower strand SEQ ID 162.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.6%; Scc
Best Local Similarity 100.0%; Pr
Matches 16; Conservative 0;
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24-NOV-2003; 2003US-00720448.
03-DEC-2003; 2003US-00727780.
14-JAN-2004; 2004US-00757803.
10-FEB-2004; 2004US-0543480P.
13-FEB-2004; 2004US-00600447.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEE32050 standard; RNA; 19 BP
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30-APR-2004; 2004WO-US013456.
24-MAY-2004; 2004WO-US016390.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-AUG-2004; 2004WO-US027366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                             targeting human ICAM1.
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The invention relates to a chemically synthesized double-stranded short interfering nucleic acid (siNA) molecule directing cleavage of interfering nucleic acid (siNA) molecule directing cleavage of confinements adhesion molecule (ICAM) RNA through RNA interference (RNAi), where each strand of the molecule is 18-23 nucleotides in length, and one strand of the molecule comprises nucleotide sequence having carficient complementarity to the ICAM RNA for the siNA molecule to direct cleavage of the ICAM RNA through RNA. The siNA is useful for downregulation or inhibition of expression of ICAM gene and for adhesion such as inflammatory diseases of the inflammatory bowel disease, atopic dermatitis and asthma, autoimmune adhesion such as inflammatory diseases e.g. rheumatoid arthritis, adhesion such as inflammatory diseases e.g. rheumatoid arthritis, cancer and proliferative diseases e.g. ovarian cancer, lung cancer, renal concer and sisease, brain injury, cerebral atrophy and congenital myopathy, respiratory diseases e.g. chronic obstructive pulmonary disease, brain injury, cerebral atrophy and congenital myopathy, respiratory diseases e.g. chronic obstructive pulmonary concertion human rown in the specification). The present sequence one strand of an siRNA through the present is the man rown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Calibration of molecular array data by employing calibration probes that
                                                                                                      Novel chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of intracellular adhesion molecule RNA through RNA interference, useful for treating diseases e.g. cancer and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for calibration of molecular array data.
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1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 19 BP; 1 A; 1 C; 1 G; 0 T; 16 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Preu. ...
                                                                                                                                                                                                   Claim 33; SEQ ID NO 162; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.6%; Score 16;
.00.0%; Pred. No.
(SIRN-) SIRNA THERAPEUTICS INC.
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Les 16; Conservative
                                    Richards I, Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     targeting human ICAM1
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                                                                        WPI; 2005-746893/76.
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molecular array. The method involves employing calibration probes that generate signals proportional to the total concentrations of labelled rarget molecules to which the molecular array probes are directed over an entire range of sample solutions and molecular arrays incorporating sets of calibration probes. Method is useful for calibrating different types of signals scanned from a molecular array, or calibrating signals scanned from different molecular arrays. The present sequence is poly (A) normalisation probe used in calibration of molecular array data The invention describes a method of predicting, diagnosing or prognosing a cardiovascular disease by detection of a polynuclectide in a biological sample comprises hybridising at least one of the polynuclectide to a nucleic acid material of a biological sample, thus forming a hybridisation complex, and detecting the hybridisation complex. The polynuclectides, polypeptides, antisense molecule, antibody and reagent are useful for preparing compositions for preventing, predicting or diagnosing, or a medicament for treating a cardiovascular disease, e.g. arteriosclerosis, isohaemia, angina pectoris, or myocardial infarction. This sequence represents a primer used to identify genes differentially generate signals proportional to total concentrations of labeled target molecules, and molecular arrays incorporating sets of calibration probes. e.g. detection Cardiovascular disease differential gene expression related primer #98 Cardiovascular disease; arteriosclerosis; ischaemia; angina pectoris; myocardial infarction; cardiant; antiarteriosclerotic; antianginal; gene therapy; differential gene expression; PCR; primer; ss. relates to a method for calibrating data scanned from a Gaps Predicting, diagnosing or prognosing a cardiovascular disease, e. angina, ischemia, myocardial infarction or arteriosclerosis by de of a polynucleotide in a biological sample comprises detecting a hybridization complex. ö Length 20; 0; Indels Seguence 20 BP; 16 A; 2 C; 0 G; 2 T; 0 U; 0 Other; 0.6%; Score 16; DB 1; Le 100.0%; Pred. No. 1.2e+03; iive 0; Mismatches 0; Schmitz G; Example 3; Page 105; 454pp; English. Disclosure, Page 14; 32pp; English Wick M, 2709 AAAAAAAAAAAAAA 2724 ВР. 02-OCT-2002; 2002WO-EP011034. 08-OCT-2001; 2001GB-00024145 1 AAAAAAAAAAAAA 16 ACA90051 standard; DNA; 20 (first entry) Local Similarity 100. nes 16; Conservative Σ Gehrmann WPI; 2003-403108/38. (FARB ) BAYER AG. WO2003031650-A2 ното варіепв 10-JUL-2003 17-APR-2003 Munnes M, ACA90051; Query Match RESULT 1506 Matches ACA90051 ð g ö Gaps

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at ftp.wipo.int/pub/published\_pct\_sequences

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
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                                                                                                                                                                                                                                            Gaps
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0
                                                                                                                                                        Score 16; DB 1; Length 20;
Pred. No. 1.2e+03;
regulated in individuals with cardiovascular disease
                                                                                Sequence 20 BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 6900; 872pp; English.
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Tang L, Shahabuddin S;
                                                                                                                                                        0.00,
100.0%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                            2076 GAAGTGACAGCTTTGA 2091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ91658 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                        4 GAAGTGACAGCTTTGA 19
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                                                                                                                                                                                                                                Matches 16; Conservative
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                                                                                                                                                                                                Best Local Similarity
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Miller S,
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                                                                                                                                                        Query Match
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung of ysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate conteainers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiathmatic, analgeaic, hypotensive, immunosuppressive and cytostatic activity, is beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administrated composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to
                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distresses syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotide containing less percentage of adenosine, targeted t
nucleic acids associated with lung airway or lung dysfunction, and
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                                         Length 20;
                                                                                 0; Indels
Sequence 20 BP; 15 A; 1 C; 0 G; 4 T; 0 U; 0 Other;
                                       Score 16; DB 1; Le
Pred. No. 1.2e+03;
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                                                                                   Mismatches
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100.001
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, Shahabuddin
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                                                                                                                                                                                                                                                                        ABD27888 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                         (first entry)
                              Query Match
Best Local Similarity 100.
Matches 16; Conservative
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, Tang L,
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reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, with a disease or condition such as pulmonary vasoconstriction, highertension, emphysema, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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.. 0.6%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 1.2e+03; cive 0; Mismatches 0; Indels 2708 TAAAAAAAAAAAAA 2723 Query Match Best Local Similarity 100.1 Matches 16; Conservative ઠે

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Gaps

TAAAAAAAAAAAA 20 유

ADH67050; RESULT 1509

Human glucocorticoid receptor-specific antisense oligonucleotide ADH67050 standard; DNA; 20 BP (first entry) 25-MAR-2004 

antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetees; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; ss; phosphorothioate backbone; 2-methoxyethyl; 2'-MOE.

Homo sapiens.

WO2003099215-A2

04-DEC-2003

20-MAY-2003; 2003WO-US016084.

20-MAY-2002; 2002US-0381857P.

(PHAA ) PHARMACIA CORP.

Crosby SD, Nalseth AE;

WPI; 2004-035034/03.

New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.

Claim 4; SEQ ID NO 3884; 985pp; English.

The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.

Seguence 20 BP; 1 A; 1 C; 2 G; 16 T; 0 U; 0 Other;

ö Gaps ö Length 20; DB 1; Le . 1.2e+03; 100.0%; Pred. ... Score 16; Pred. No. 2709 AAAAAAAAAAAAA 2724 20 AAAAAAAAAAAAAA Conservative Best\_Local Similarity Matches 16; Conserv Query Match

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ADK76466 standard; DNA; 20

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ADX76466;

20-MAY-2004 (first entry)

Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.

Chimeric phosphorothioate oligonucleotide to target Nav1.3 #3800.

Synthetic.

WO2004016754-A2.

26-FEB-2004.

14-AUG-2003; 2003WO-US025465.

14-AUG-2002; 2002US-0403416P.

(PHAA ) PHARMACIA CORP

Roberds SL;

WPI; 2004-203785/19.

New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.

Claim 4; SEQ ID NO 3800; 417pp; English.

The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represente a chimeric phosphorothicate oligonacleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.

Sequence 20 BP; 1 A; 3 C; 0 G; 16 T; 0 U; 0 Other;

Gaps ö 0.6%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 1.2e+03; ive 0; Mismatches 0; Indels Query Match 0.6 Best Local Similarity 100. Matches 16; Conservative

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WO2005118843-A1
                                     Homo sapiens.
                                                                                                                                                                                                                                                              (DRD2) gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                            Included and declare encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3. e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, pain fine neuropathy, trigeminal neuropathy, arthritic pain, lower back pain, pain from burns, migraine headachy, archritic pain, acute pain, pain from burns, migraine headach a childhood seizure disorder such as childhood seizure disorder including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with Wilman Navi.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                      New antisense compound targeted to a nucleic acid molecule encoding May1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                               Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human dopamine receptor D2 (DRD2) DNA oligonucleotide SEQ ID NO:2629.
                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to an antisense compound targeted to
                                                                                           Chimeric phosphorothioate oligonucleotide to target Navl.3 #2548.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 1.2e+03; ative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 2548; 417pp; English.
                        BP.
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                     ADK75214 standard; DNA; 20
                                                                    (first entry)
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Matches 16; Conservative
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                                                                    20-MAY-2004
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                                                                                                                                                               Synthetic.
                                             ADK75214;
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RESULT 1511
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The invention relates to a method of identifying a genetic profile associated with a neurological, psychiatric or psychological condition, phenotype or state including a sub-threshold neurological, psychiatric or psychological condition, phenotype or state in an individual, comprising screening individuals for a polymorphism in a genetic locus comprising companies receptor D2 (DRD2) gene. The invention also relates to a genetic mutation providing a genetic marker for a neurological, psychiatric, or psychological condition, state or phenotype in an individual, where the presence of a 957 polymorphism is indicative of a psychiatric or psychological condition, phenotype or state. The compositions and methods are useful condition, phenotype or state. The compositions and methods are useful condition, phenotype or state. The compositions and methods are useful condition a genetic profile associated with a neurological, psychiatric or psychological condition. The method enables clinicians to make a genetic-based diagnosis of a neurological, psychiatric or psychological condition and can thereby implement treatment or correct preventative or symptom-ameliorating protocols to reduce the adverse consequences of the condition. This sequence represents a human dopamine incention is the scope of the condition.
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cationic; gene therapy; rheumatoid arthritis; targetting;
distal administration; delivery; macrophage; uptake; polyamine; PCR;
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for a
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Diagnosis; therapeutic; neurological disease; psychiatric disorder; neuropsychologic disorder; dopamine receptor D2; DRD2; ss.
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100.08; Pre-
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Matches 16; Conserv
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The present invention relates to human acetyl-Coenzyme A-carboxylass-
alpha (ACC-alpha; see ABZ79442), which can be used for in vitro diagnosis
of cancer (or of an increased risk of developing it), by detecting ACC-
alpha gene mutations or polymorphisms, or altered ACC-alpha protein
expression, relative to a control population. The method is particularly
used to diagnose cancer, especially of bresst or overy, or for assessing
the risk of developing such cancers. The present sequence is a PCR
primer, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                               In vitro diagnosis of cancer, particularly breast and ovarian cancer, or susceptibility, comprises detecting alterations in the acetyl coenzyme A-carboxylase alpha gene or protein expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; polycomb group protein; EZH2; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy;
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                                                                                                                                                                                                                                                                                                    Sinilnikova-Erard
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Pred. No. 1.2e+03;
0; Mismatches 2;
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11-MAR-2002; 2002US-0365124P.
06-UUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                    Magnard CM,
                                                                                                                                                  2001FR-00007740.
2002FR-00002788.
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                                                                                                12-JUN-2002; 2002WO-FR002015
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Matches 17; Conservative
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WO2002100896-A2
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05-MAR-2002;
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                                                 19-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transcription or KANA and DNA encoding numan IL-10 in multine tissue amplified DNA was then detected via in situ hybridisation with a human IL-10 expression vector. The amplified DNA was then detected via in situ hybridisation with a human IL-10 use of a derivatised cationic liposome containing an agent of interest (e.g., an IL-10 expression vector) to deliver the agent to a site that is liposome administration. The derivatised cationic liposome comprises a cationic liposome forming entity (e.g., a lipid), and a head-group and/or ligand which increases macrophage uptake of the cationic liposome. Such liposome can be used for the delivery of a cationic liposome. Such liposomes can be used for the delivery of a cationic liposome. Such liposome macrophage uptake is used coin miprove therapeutic agent, and is especially useful for the amelioration of cationic liposome macrophage uptake is used to improve targetting and thereby make more efficient use of the cationic liposome and agent of interest If cholesterol is used as the cationic liposome cationic liposome-forming entity is linked to the head group via a cationic liposome-forming entity is linked to the head group via a cationic liposome-forming entity is linked to the head group via a cationic liposome-forming entity is linked to the head group via a cationic liposome, and also increases the overall positive charge on the liposome. As polyamines occur naturally in cells, correctly considered by a binding and cationic by the liposome. As polyamines occur naturally in cells,
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                                                                                                                                                                                                                                                                                                                                                                                                       Use of composition comprising combination of derivatized cationic liposomes and active agent in gene therapy, especially for amelioration of established arthritis by IL-10 gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a human IL-10 (interleukin-10) PCR primer #1, used with primer #2 (AAZ32680) to amplify cDNA (generated via reverse transcription of RNA) and DNA encoding human IL-10 in murine tissue previously transfected with a human IL-10 expression vector. The
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                                                                                                                                                                                                                                                                                                         Ley SV;
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                                                                                                                                                                                                                                                                                                         Duffels AT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 8 A; 6 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                         WOO P,
                                                                                                                                                                                                                                                       (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
                                                                                                                                                                                                                                                                                                         Etheridge CJ, Fellowes R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 30; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1542 TGAAAACCAAGACCCGGAC 1560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGAGAACCAAGACCCAGAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABZ79441 standard; DNA; 19 BP
                                                                                                                                                    99WO-GB001171
                                                                                                                                                                                                      98GB-00008268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-633970/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast; ovary;
       Homo sapiens
                                                      WO9954344-A1
                                                                                                                                                       16-APR-1999;
                                                                                                                                                                                                        17-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-2003
                                                                                                                                                                                                                                                                                                         Miller AD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ79441;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1514
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Gaps

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Length 19; Indels 88.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human polycomb group protein EZH2 gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense of soligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically motified, can contain deoxyribonucleotides, and can be chemically synthesised. expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in synthesised. The siNAs are used to modulate expression of the EZH2 convict of siNA. The siNAs are used to modulate expression of the EZH2 gene in cells, tissue explants or organisms (e.g., by ex vivo gene the capture of a grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation canneric enganger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human EZH2 targeted
                                                                                                                                                                                 New short interfering nucleic acid, useful e.g. for treatment and diagnosis of prostatic cancer, downregulates expression of the EZH2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; polycomb group protein; EZH2; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; cancer; restenosis; drug screening; diagnosis; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human EZH2 transcript target sequence/siNA upper strand, SEQ ID 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.8; DB 1; Length 1
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 1 C; 1 G; 0 T; 15 U; 0 Other;
                                                                                                          Usman N;
                                                                                                      Beigelman L, Haeberli P,
                                                                                                                                                                                                                                              Example 7; Page 121; 140pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2706 ACTAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 ACTTGAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF92943 standard; RNA; 19 BP
  19-NOV-2002; 2002US-0427467P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-2003; 2003WO-US004402.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2002; 2002US-0358580P.
                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double stranded siNA.
                                                                                                                                            WPI; 2003-712612/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003070887-A2
                                                                                                    Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF92943;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 1516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human polycomb group protein EZH2 gene by CC RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense coligonucleotide and an antisense oligonucleotide. Specifically, the siNAs coligonucleotide short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kite for the in vitro or in synthesised. The invention also relates to kite for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the EZH2 conditions. They may be used for treating cancer. The siNAs are also conditions. They may be used for treating cancer. The siNAs are also conditions. They may be used for treating cancer. The siNAs are also and validation, genetic engineering, pharmacogenomics, studying gene condition, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human EZH2 targeted condition.
                                                                                                                                                                                                                                                                             New short interfering nucleic acid, useful e.g. for treatment and diagnosis of prostatic cancer, downregulates expression of the EZH2 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; eytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; exbB2; c-erb-B-2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 15.8; DB 1; Length 19; 34.2%; Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human HER2 (EGFR2) siNA lower strand, SEQ ID NO:496.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 15 A; 1 C; 1 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                    Usman N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Mismatches
                                                                                                                                                                                                    Mcswiggen J, Beigelman L, Haeberli P,
                                                                                                                                                                                                                                                                                                                                       Example 7; Page 121; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2706 ACTAAAAAAAAAAAAAA 2724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ACUUGAAAAAAAAAAA 19
                  2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2002US-0427467P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL79331 standard; RNA; 19 BP
                                                                                                 19-NOV-2002; 2002US-0427467P
15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84.2%;
                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 16; Conservat
                                                                                                                                                                                                                                       WPI; 2003-712612/67.
                  06-JUN-2002;
29-AUG-2002;
                                                            05-SEP-2002;
                                                                              09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL79331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              seguence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1517
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WO2003070912-A2

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The invention relates to short interfering nucleic acids (siNh) which downregulate expression of one or more human epidermal growth factor receptor (ESPR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNNs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vector to the in vivo delivery of siNA; conjugates and or complexes of siNA; and vectors that express sinA. The siNAs are used for or a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The siNAs are also useful for drug screening, diagnosis, therapeutic target dentification and validation, genetic engineering, pharmacogennics, studying gene function, and gene mapping (e.g., of single nucleotide and termand of a partey. EEER2 (EGFR2)-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                              Jamison S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 16 A; 2 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                              Fosnaugh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 496; 171pp; English.
                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Pavco P, Beigelman L,
                                                                                                                                                                                                                                         05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408239P.
18-SEP-2002; 2002US-0251117.
21-OCT-2002; 2002US-00277494.
                                                                                                                             2002US-0363124P.
2002WO-US016840.
2002US-00163552.
                                                                                                                                                                                                       2002US-0393924P
2002US-0406784P
                                                                      20-FEB-2003; 2003WO-US005045
                                                                                                                                                                                    2002US-0386782P
                                                                                                                                                                                                                                                                                                                   15-JAN-2003; 2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 84.2%;
                                                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-697612/66.
                                                                                                                                                                                  06-JUN-2002;
03-JUL-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                             29-MAY-2002;
06-JUN-2002;
                                 28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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ö Gaps ö 0.6%; Score 15.8; DB 1; Length 19; 4.2%; Pred. No. 1.2e+03; 2708 TAAAAAAAAAAAAAA 2726 1 UAAAAAACAAAACAAAA 19 용

ADL79082 standard; RNA; 19 BP (first entry) 20-MAY-2004 ADL79082; RESULT 1518

Human HER2 (EGFR2) transcript target sequence/siNA upper strand, SEQ:247 RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mikNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; exbB2; c-erb-B-2; target sequence; ss. 2002US-0363124P. 2002WO-US016840. 2002US-00163552. 2002US-0386782P. 20-FEB-2003; 2003WO-US005045. 03-JUL-2002; 2002US-0393924P 29-AUG-2002; 2002US-0406784P 05-SEP-2002; 2002US-0408378P 2002US-0408378P. 2002US-0409293P. 19-SEP-2002; 2002US-00251117. 21-OCT-2002; 2002US-00277494. 15-JAN-2003; 2003US-0440129P. 2002US-0358580P WO2003070912-A2. 29-MAY-2002; 06-JUN-2002; 06-JUN-2002; Homo sapiens. 09-SEP-2002; 20-FEB-2002; 11-MAR-2002; 28-AUG-2003 

Jamison S; Beigelman L, Fosnaugh K, Pavco P, Mcswiggen J,

WPI; 2003-697612/66.

(RIBO-) RIBOZYME PHARM INC.

New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth factor receptor gene.

Example 3; SEQ ID NO 247; 171pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor receptors (EGFR) genes (including HRR1, HRR2 HRR3 and HRR4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and matisense oligonucleotide. Specifically, the siNAs include and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short cinterfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Cc an contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also crace contain deoxyribonucleotides, and can be chemically synthesised. The invention also can contain deoxyribonucleotides, and can be chemically synthesised. The invention also and/or complexes of siNA; and vectors that express siNAs. The siNAs are cused to modulate expression of EGFR genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants or creating a wider range of cancers such as breast and ovarian cancer. The treatment of a variety of conditions. They may be used for treating a wider range of cancers such as breast and ovarian cancer. The cidentification and validation, genetic engineering, pharmacogenomics, cidentification and validation, and gene mapping (e.g., of single nucleotide bluman HER2 (EGFR2)-targeted double-stranded siNA, which is identical to the here the HER2 transcript target sequence.

Sequence 19 BP; 1 A; 0 C; 2 G; 0 T; 16 U; 0 Other;

Gaps ö Query Match 0.6%; Score 15.8; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 1.2e+03; Matches 17; Conservative 0; Mismatches 2; Indels

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Protein tyrosine phosphatase siRNA sequence, SEQ ID No 197.
                                                                                                                                                                                                                            Example 3; SEQ ID NO 197; 234pp; English.
TAAAAAAAAAAAAAA 2726
          TAAAAAACAAAAA 1
                                     ΒP
                                                                                                                                                  23-MAY-2002; 2002US-0383249P.
14-APR-2003; 2003US-0462942P.
                                                                                                                                        23-MAY-2003; 2003WO-US016651
                                   ADH01585 standard; RNA; 19
                                                        11-MAR-2004 (first entry)
                                                                                                                                                                             Lewis SP, Klinghoffer R,
                                                                                                                                                                                        WPI; 2004-035036/03.
                                                                                                                                                                  (CEPT-) CEPTYR INC.
                                                                                                                                                                                                                  diabetes or cancer
                                                                                                                   WO2003099227-A2.
                                                                                                         Unidentified
                                                                                                                              04-DEC-2003.
2708
                                              ADH01585;
          13
                         RESULT 1519
                                ADH01585
          셤
                                         ઠે
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Gaps ö 0.6%; Score 15.8; DB 1; Length 19; 78.9%; Pred. No. 1.2e+03; 2; Indels Sequence 19 BP; 6 A; 4 C; 7 G; 0 T; 2 U; 0 Other; 2; Mismatches Best Local Similarity 78.9 Matches 15; Conservative Query Match

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gene typing; polymorphic microsatellite loci; PML; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; cow; microsatellite; PCR; primer; 88. Cow prion protein microsatellite locus primer #69. 2436 TGAAGAAGCAGGAGCTGCT 2454 Example 3; Page 28; 64pp; German. Han Y; 1 ugaagaagcagcagcu 19 ADO81057 standard; DNA; 19 BP 2002DE-01036711. 09-AUG-2002; 2002DE-01036711. (first entry) s, Conservative (UYHO-) UNIV HOHENHEIM. Geldermann H, Preuss WPI; 2004-215730/21. length of amplicons. Local Similarity DE10236711-A1. 09-AUG-2002; 29-JUL-2004 26-FEB-2004. 17; Bos taurus. ADO81057; Query Match RESULT 1520 Matches ADO81057, ઠ The invention relates to a novel isolated small interfering RNA (SIRNA)

polynucleotide, comprising at least one nucleotide sequence from any of
the 20 fully defined sequences given in the specification. The invention
further relates to: a pharmaccutical composition comprising a new siRNA

construct, comprising a polynucleotide that is capable of directing
cranscription of an siRNA; a host cell transformed or transfected with

the above recombinant nucleic acid construct; a method for interfering

the above recombinant nucleic acid construct; a method for interfering

with expression of a protein tyrosine phosphatase (PTP) 1B polypeptide, or

the variant; a method for indentifying a component of a PTP1B signal

transduction pathway; a method for indentifying a component of a PTP1B signal

transduction state in a cell; a method for altering a Jak2 protein

phosphorylation state in a cell; and a method for treating a Jak2 protein

construct an orectic, antiinflammatory, neuroprotective, cytostatic,

antidiabetic, anorectic, antiinflammatory, neuroprotective, cytostatic,

immunosuppressive, and antimicrobial. The novel siRNA polynucleotides can

be used in gene therapy to treat disorders. The composition and methods

cransduction, such as diabetes, obesity, hyperglycaemia-induced

transduction, apoptosis, inflammation, neurodespensative disorders, cancer, autoimmune diseases or infection. This polynuclectide sequence represents an siRNA phosphatase of the invention. owdail incertering RNA; siRNA; protein tyrosine phosphatase; PTP; PTP1B; insulin receptor protein phosphorylation; Jak2; antidiabetic; anorectic; antiinflammatory; neuroprotective; cytostatic; immunosuppressive; antimicrobial; gene therapy; ss; siRNA. New small interfering polynucleotide that modulates protein tyrosine phosphatase (PTP)1B polypeptide signal transduction, useful for treating disorders associated with altered PTP1B signal transduction, e.g.

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The invention describes a method of typing (M1) a gene (1) that has one or more polymorphic microsatellite loci (PML). The method comprises: PCR amplification of at least one DNA region of (1) that includes PML, using as template a DNA sample containing at least one segment of (1); and catermining the length of the resulting amplicon(s). Also described are: canethod of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or concerp. The method is used to identify microsatellite markers, in a disease-related gene, that are associated with a predisposition to diseases and for predisgnosis of such diseases, especially prion diseases of diseases and for predisgnosis of such diseases, especially prion diseases concerp. This predisposition to metabolic diseases, also to type genes that encode milk proteins, mailgnant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins.

This processes is also to type genes that encode milk proteins.

Described the processes of the processes of the processes of the processes of the processes and for predisposition for the processes of the processes and for predisposition for the processes of the processes and processes are processes. The method is simpler, quicker and particularly less expensive than known methods based on sequencing.
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Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.
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89.5%; Pred. No. 1.2e+03;
tive 0; Mismatches 2; Indels
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The invention describes a RNA interference (iRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequences have cone or more asymmetrical 2'-O alkyl modifications, the antisense sequence asymmetrical phosphorothioate modifications and the antisense sequence targets a human gene sequence. Also described are: a pharmaceutical preparation comprising (I); reducing (M1) apoB-100 contevels or glucose-6-phosphatase levels in a subject; producing (I); catabilising (I), involves selecting a sequence with activity and introducing one or more asymmetrical modification in the sequence, where the modification decreases nuclease sensitivity while not decreasing its activity; a kit comprising (I) and instruction for its use; and a device that can be dispense or administer a composition comprising (I). (I) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. (M1) is useful for reducing apoB-100 levels or glucose-6-phosphatase levels. The subject is suffering from a disorder characterised by elevated or otherwise unwanted expression of apoB-100, elevated or otherwise unwanted levels and or alipid metabolism. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Interference RNA agent useful for treating dyslipidemias, coronary artery disease, diabetes, cancer or neurological disease, comprises sense sequence and antisense sequence which has specific modifications.
                                                                                                                                                                                                                                                                                                  cytostatic; anticonvulsant; nootropic; muscula; anti-HIV; RNA interference; IRNA, antisense technology; lipid metabolism; cholesterol imbalance; dyslidemia hypercholesterolaemia; coronary artery disease; CAD; coronary heart disease; CHD; atherosclerosis; hepatic glucose production; albetees; cancer; breast cancer; glucose-metabolism-related disorder; diabetees; cancer; breast cancer; colon cancer; lung cancer; neurological disease; Hultington disease; spinocerebellar ataxia; viral disease; AIDS; glucose-6-phosphatase; sg.
                                                                                                                                                                                                                                                                  antilipemic; cardiant; vasotropic; antiarteriosclerotic; antidiabetic;
                                                                                                                                                                                                   Human glucose-6-phosphatase oligonucleotide seqid 5367.
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2003US-0454265P.

2003US-0455050P.

2003US-0462894P.

2003US-046372P.

2003US-0463612P.

2003US-0469612P.

2003US-0469612P.

2003US-0494597P.

2003US-0494597P.

2003US-0404697P.

2003US-0404697P.

2003US-0404697P.

2003US-0404697P.
ADR80868 standard; DNA; 19 BP.
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                                                                                                                                    (first entry)
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interference RNA; iRNA; cholesterol moiety; apoB; glucose-6-phosphatase;
lipid metabolism; cholesterol imbalance; dyslipidaemia;
familial combined hyperlipidaemia; acquired hyperlipidaemia;
disorder is chosen from the HDL/LDL cholesterol imbalance, dyslipidaemias, hypercholestorolaemia, statin-resistant hypercholestorolaemia, coronary artery disease (CAD), coronary heart disease (CAD), and atherosclerosis. (I) is administered to subject to inhibit hepatic glucose production or for treating glucose-metabolism-related disorder e.g. diabetes or type-2 diabetes. (I) is useful for treating the diseases as mentioned above, cancer (e.g. breast, colon or lung cancer), neurological disease (e.g., Huntington disease or spinocerebellar ataxia) or viral disease (e.g., AIDS). This sequence the represents a human glucose-6 phosphates entisense oligonucleotide that
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glucose-6-phosphatase antisense inhibition target seqid 5367.
                                                                                                                                                                               can be used to control glucose-6-phosphatase gene expression.
                                                                                                                                                                                                                                                Length 19;
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Pred. No. 1.2e+03;
0; Mismatches 2;
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14-APR-2003; 2003US-0462894P.
17-APR-2003; 2003US-046565P.
25-APR-2003; 2003US-0465665P.
25-APR-2003; 2003US-0465602P.
09-MAY-2003; 2003US-0465602P.
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11-AUG-2003; 2003US-0494597P.
26-SEP-2003; 2003US-0506341P.
09-OCT-2003; 2003US-0510246P.
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                                                                                                                                                                                                                                                                                   17; Conservative
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Best Local Similarity
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The invention describes an interference RNA (IRNA) agent (I) comprising a sense sequence and an antisense sequence, where the sense sequence comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence comprises one or more cholesterol moieties, and the antisense sequence cargets a human gene sequence. The following are disclosed: a pharmaceutical composition comprising (I); and a device for administering (I) into a patient. (I) is useful for reducing apollo levels or glucose of phosphatase levels in a subject. (I) targets a sequence identical to any one of sequences as given in the specification. (I) comprises a cholesterol moiety is coupled to a sense strand. (I) further comprises a second cholesterol moiety. The second cholesterol complety is coupled to a sense strand. (I) has 21 or more nucleotides. The duplex region of (I) is 19 nucleotides in length. The subject is suffering from a disorder having elevated or otherwise unwanted levels of expression of apo-B-100, elevated or otherwise unwanted levels of cholesterol, and/or disregulation of lipid metabolism. The disorder is combined hyperlipidaemia or acquired hyperlipidaemia, coronary company of the service of t
New interference RNA agent comprising sense sequence and antisense sequence having cholesterol moleties, useful for reducing apoB-100 levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       artery disease, coronary heart disease and atherosclerosis, preferably statin-resistant hypercholesterolaemia. (1) is administered to a subject to inhibit hepatic glucose production or for treating glucose-metabolism-related disorders e.g., type-2 diabetes or glitaxzone-resistant diabetes. (1) has endonuclease or exonuclease resistance. This sequence represents a human glucose-6-phosphatase pallindromic sequence that may be useful as
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X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi;
cytostatic; cancer; gene silencing.
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89.5%; Pred. No. 1.2e+03;
trive 0; Mismatches 2; Indels
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                                                                                                          Example 4; SEQ ID NO 5367; 324pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target for antisense inhibition.
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                                                             or glucose-6-phosphatase levels.
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; 2003US-00693059.
; 2003US-00720448.
; 2004US-0075780.
; 2004US-057800.
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; 2004US-00760447.
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les 17; Conservative
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10-FEB-2004;
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This invention describes novel chemically synthesized double stranded short interfering nucleic acid (siRNA) molecules which direct cleavage of a X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference (RNAi), where each strand of the siRNA molecule is about 18-23 nucleotides in length and one strand of the siRNA molecule comprises nucleotides in length and one strand of the siRNA molecule comprises.

CC nucleotides in length and one strand of the siRNA molecule comprision comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecule. The sense and comprising the sense region are 2'-deoxy purine cucleotides in the sense region are 2'-deoxy purine cucleotides in the sense region are 2'-deoxy purine cucleotides and the pyrimidine nucleotides are 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region includes a terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. The terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. The purine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides of the cantisense region comprises a glyceryl modification at a 3'-end of the antisense region. The antisense region comprises a glyceryl modification at a 3'-end of the antisense region. About 19 nucleotides of each fragment of the siRNA molecule are base-paired to the complementary nucleotides of the other CC antisense region optionally includes a phosphotosphate group. The XIAP RNA comprises Genbank Accession No. NM 001167. The chemically synthesized comprises dowble stranded short interfering nucleic acid elsaperium for the remained region optionally includes a phosphate group. The XIAP RNA comparing a comparing
                                                                                                                                                                                                             New chemically synthesized double stranded short interfering nucleic acid that directs cleavage of an X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference, useful in preparing a composition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ds; primer; short interfering RNA; siRNA;
X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi;
cytostatic; cancer; gene silencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in preparing a composition for treating cancer. ADX86130-ADX87180 represent siRNA molecules which are used in RNA interference mediated
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                                                                                                                                                                                                                                                                                                                                      Claim 33; SEQ ID NO 372; 202pp; English.
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                                                                    (SIRN-) SIRNA THERAPEUTICS INC.
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30-APR-2004; 2004WO-US013456.
24-MAY-2004; 2004WO-US016390.
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                                                                                                                  Mcswiggen J, Chowrira
                                                                                                                                                                  WPI; 2005-163247/17.
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ADX86332/c
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16-APR-2004;

ADX86799 standard; RNA; 19 BP

RESULT 1525 4DX86799

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This invention describes novel chemically synthesized double stranded short interfering nucleic acid (siRNA) molecules which direct cleavage of a X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference (RNAi), where each strand of the siRNA molecule is about 18-23 nucleotides in length and one strand of the siRNA molecule comprises nucleotide sequence having sufficient complementarity to the XIAP RNA. The siRNA molecules can be used to make a cytostatic composition comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecule. The pyrimidine nucleotides in the sense region are 2'-deoxy purine nucleotides and the pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides. The fragment comprising the sense region includes a terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. The terminal cap moiety is an inverted deoxy abasic moiety. The pyrimidine nucleotides of the antisense region comprises a phosphorothioat comprise cutients are 2'-deoxy-2'-fluoro pyrimidine nucleotides are 2'-deoxy-2'-fluoro comprises a phosphorothioat centre antisense region comprises a phosphorothioat end comprises a glyceryl modification at a 3' end of the antisense region comprises a glyceryl modification at a 3' end of the antisense region comprises a glyceryl modification at a 3' end of the changes antisense region comprises of each fragment of the bithat
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in preparing a composition for treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SIRN-) SIRNA THERAPEUTICS INC.
                                                                                                                                                                                      2003US-0493561P.
2003US-00693059.
2003US-00720448.
2003US-00727780.
2004US-00757803.
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2004US-00780447.
2004US-00826966.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating cancer.
                                             WO2005014811-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J,
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03-DEC-2003;
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                                                                                            17-FEB-2005
  Synthetic.
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represent sirna molecules which are used in RNA interference mediated Sequence 19 BP; 4 A; 0 C; 2 G; 0 T; 13 U; 0 Other; Inhibition of XIAP gene expression. 0.68;

cancer.

ö Gaps ö Score 15.8; DB 1; Length 19; Pred. No. 1.2e+03; 0; Mismatches 2; Indels 89.5%; Query Match
Best Local Similarity 89.5'
Matches 17; Conservative

2703 TGTACTAAAAAAAAAA 2721

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TCTACTAAAAATAAAAAA 1

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New chemically synthesized double stranded short interfering nucleic acid that directs cleavage of an X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference, useful in preparing a composition for ds; primer; short interfering RNA; siRNA; X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi; cytostatic; cancer; gene silencing. Claim 33; SEQ ID NO 670; 202pp; English. XIAP targeting siRNA SEQ ID NO 670. (SIRN-) SIRNA THERAPEUTICS INC. 2003US-00693059 2003US-00720448 2003US-00727780 2004US-00757803. 2004US-0543480P. 2004US-00826966 2004WO-US013456 24-MAY-2004; 2004WO-US016390 06-AUG-2004; 2004WO-US025589 2003US-0493561P 2004US-00780447 Chowrira BM; 05-MAY-2005 (first entry) WPI; 2005-163247/17. WO2005014811-A2. treating cancer Mcswiggen J, 08-AUG-2003; 16-APR-2004; 03-DEC-2003; 14-JAN-2004; 13-FEB-2004; 23-OCT-2003; 24-NOV-2003; 17-FEB-2005. Synthetic. ADX86799; 

This invention describes novel chemically synthesized double stranded short interfering nucleic acid (siRNA) molecules which direct cleavage of a X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference (RNAi), where each strand of the siRNA molecule is about 18-23 nucleotides in length and one strand of the siRNA molecule comprises nucleotide sequence having sufficient complementarity to the XIAP RNA. The siRNA molecules can be used to make a cytostatic composition comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecule. The pyrimidine composition comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecules. The pyrimidine nucleotides in the sense region are 2'-deoxy purine conclectides and the pyrimidine nucleotides are 2'-deoxy purine conclectides and the pyrimidine nucleotides are 2'-deoxy purine conclection of the fragment comprising the sense region. The terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. The terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprises a phosphorothicate contineers region are 2'-deoxy-2'-fluoro pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides of the antisense region. The antisense region comprises a glyceryl modification at a 3'-deoxy purine cucleotides present in the antisense region comprises a phosphorothicate contineers region comprises a glyceryl modification at a 3'-deoxy purine antisense region. About 19 nucleotides of each fragment of the siRNA molecule are base-paired to the complementary nucleotides of the other fragment of the siRNA. The 5'-end of the fragment comprising the sendent of the siRNA molecule are base-paired to the complementary nucleotides of the other fragment of the siRNA molecule are done of the complementary nucleotides.

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antisense region optionally includes a phosphate group. The XIAP RNA comprises Genbank Accession No. NM 001167. The chemically synthesized double stranded short interfering Tucleic acid (siRNA) molecule is useful in preparing a composition for treating cancer. ADX86130-ADX87180 represent siRNA molecules which are used in RNA interference mediated inhibition of XIAP gene expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ds; primer; short interfering RNA; siRNA;
X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi;
cytostatic; cancer; gene silencing.
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                                                                                                                                                  0.6%; Score 15.8; DB 1; Length 19; 73.7%; Pred. No. 1.2e+03; tive 3; Mismatches 2; Indel8
                                                                                                                    Sequence 19 BP; 13 A; 2 C; 0 G; 0 T; 4 U; 0 Other;
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2003US-00720448.
2003US-00727780.
2004US-00757803.
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2004US-00780447.
2004US-00826966.
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                                                                                                                                           Query Match
Best Local Similarity 73.7
Matches 14; Conservative
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10-FEB-2004;
13-FEB-2004;
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24-NOV-2003;
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The purine nucleotides in the sense region are 2'-deoxy purine nucleotides and the portining nucleotides in the sense region are 2'-deoxy purine nucleotides and the pyrimidine nucleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides. The fragment comprising the sense region includes a terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region. The terminal cap moiety is an inverted deoxy absaic moiety. The pyrimidine nucleotides of the antisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides of the concleotides are 2'-O-methyl purine nucleotides. The purine nucleotides are 2'-O-methyl purine nucleotides. The antisense region comprises a phosphorothicate internucleotide present in the antisense region comprises a phosphorothicate concleotides. The antisense region comprises a glyceryl modification at a 3'-end of the antisense region. About 19 nucleotides of each fragment of the siRNA comprises a glyceryl modification to the siRNA comprises genbank Accession No. Nucleotides a phosphate stranded short interfering nucleic acid (siRNA) molecule is useful in preparing a composition for treating cancer. ADX86130-ADX87180 comprises the siRNA molecules which are used in RNA interference mediated inhibition of XIAP gene expression.
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 nucleotides in the sense region are 2'-O-methyl pyrimidine nucleotides.
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X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi;
cytostatic; cancer; gene silencing.
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24-NOV-2003; 2003US-00720448.
03-DEC-2003; 2003US-0072780.
14-JAN-2004; 2004US-00757803.
10-FBB-2004; 2004US-0543480P.
13-FBB-2004; 2004US-0048447.
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les 17; Conservative
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Matches
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This invention describes novel chemically synthesized double stranded short interfering nucleic acid (siRNA) molecules which direct cleavage of a X-linked inhibitor of apoptoais protein (XIAP) RNA via RNA interference (RNAi), where aech strand of the siRNA molecule is about 18-23 nucleotides in length and one strand of the siRNA molecule comprises nucleotide sequence having sufficient complementarity to the XIAP RNA. The siRNA molecules can be used to make a cytostatic composition comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecule. The pyrimidine nucleotides in the sense region are 2'-deoxy purine concleotides in the sense region are 2'-deoxy purine concleotides and the pyrimidine nucleotides are 2'-deoxy purine concleotides and the pyrimidine nucleotides are 2'-deoxy purine comprising the sense region includes at terminal cap moiety at a 5'-end, a 3'-end, or both of the 5' and 3' ends of the fragment comprising the sense region includes can tisense region are 2'-deoxy-2'-fluoro pyrimidine nucleotides are 2'-deoxy purine concleotides are 2'-deoxy-2'-fluoro pyrimidine nucleotides are 2'-deoxy-2'-fluoro contrises a phosphorothioat The arrisense region comprises a phosphorothioat The contriber antisense region comprises a phosphorothioat The arrise antisense region comprises a glyceryl modification at a 3' end of the contriber antisense region comprises a glyceryl modification at a 3' end of the contriber antisense region of each fragment of the strand of strand 
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New chemically synthesized double stranded short interfering nucleic acid that directs cleavage of an X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference, useful in preparing a composition for
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X-linked inhibitor of apoptosis protein; XIAP; RNA interference; RNAi;
cytostatic; cancer; gene silencing.
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                                                                                                                                               Claim 33; SEQ ID NO 296; 202pp; English.
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                                                                                             treating cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-MAY-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1528
     윱
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This invention describes novel chemically synthesized double stranded short interfering nucleic acid (siRNA) molecules which direct cleavage of short interfering nucleic acid (siRNA) molecules which direct cleavage of a X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference (RNAi), where each strand of the siRNA molecule is about 18-23 nucleotides in length and one strand of the siRNA molecule comprises on be used to make a cytostatic composition comprising the siRNA molecule in a carrier or diluent. The sense and antisense strands are connected via a linker molecule. The pyrimidine comprising the sense region are 2'-deoxy purine nucleotides in the sense region are 2'-deoxy purine nucleotides and the pyrimidine nucleotides are 2'-deoxy purine nucleotides. The fragment comprising the sense region includes or the fragment comprising the sense region includes care the fragment comprising the sense region includes can inverted deoxy abasic moiety. The pyrimidine nucleotides of the continent are 2'-deoxy-2'-fluoro pyrimidine nucleotides are 2'-deoxy-2'-fluoro comprises a phosphorothioar. The antisense region comprises a glyceryl modification at a 3' end of the antisense region are 2'-deoxy-purine nucleotides are 2'-deoxy-pur
                                                                                                                                                                                                                                                                                                                                                                                                          New chemically synthesized double stranded short interfering nucleic acid that directs cleavage of an X-linked inhibitor of apoptosis protein (XIAP) RNA via RNA interference, useful in preparing a composition for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 molecule are base-paired to the complementary nucleotides of the other fragment of the siRNA. The 5'-end of the fragment comprising the antisense region optionally includes a phosphate group. The XIAP RNA comprises Genbank Accession No. NM 001167. The chemically synthesized double stranded short interfering nucleic acid (siRNA) molecule is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in preparing a composition for treating cancer. ADX86130-ADX87180 represent siRNA molecules which are used in RNA interference mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gape
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Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 1 A; 1 C; 0 G; 0 T; 17 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 33; SEQ ID NO 763; 202pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inhibition of XIAP gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2709 AAAAAAAAAAAAAAAA 2727
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                                           03-DEC-2003; 2003US-00727780.
14-CAN-2004; 2004US-0054803.
10-FEB-2004; 2004US-0543480P-
13-FEB-2004; 2004US-0080447.
16-APR-2004; 2004WG-US01656.
24-MAY-2004; 2004WG-US011456.
                                                                                                                                                                                                                                                            (SIRN-) SIRNA THERAPEUTICS INC.
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                                                                                                                                                                                                                                                                                                             Chowrira BM;
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Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating cancer.
                                                                                                                                                                                                                                                                                                                Mcswiggen J,
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Tue Nov

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interference; gene silencing; short interfering RNA; siRNA; cancer;
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Human IL-4R transcript target sequence/siRNA sense strand, SEQ ID:389.
                                                                                                                                                   hyperproliferation; neoplasm; cytostatic; viral infection; infection; virucide; inflammation; antiinflammatory; autoimmune disease;
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Homo sapiens.

US2005143333-A1.

30-JUN-2005

09-JUN-2004; 2004US-00863973

2002WO-US015876. 2002US-0386782P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 2003US-0440129P. 2001US-0311865P. 2002US-0358580P. 2002US-0362016P. 2002US-0363124P. 06-MAR-2002; 11-MAR-2002; 20-MAY-2002; 06-JUN-2002; 09-SEP-2002; 39-AUG-2002; 05-SEP-2002;

2003US-00693059. 2003WO-US004566 2003WO-US005028 2003WO-US005346 2003US-00427160 2003US-00444853 2003US-00727780 2004US-00757803 23-OCT-2003; 24-NOV-2003; 03-DEC-2003; 15-JAN-2003; 20-FEB-2003; 20-FEB-2003; 14-JAN-2004; 23-MAY-2003;

2004WO-US016390. 2004US-00826966 24-MAY-2004; 30-APR-2004 16-APR-2004

2004US-0543480P 2004US-00780447 (SIRN-) SIRNA THERAPEUTICS INC.

Richards I, Polisky B, Mcswiggen J;

WPI; 2005-457799/46.

Novel chemically synthesized double stranded short interfering nucleic acid molecule useful for cleaving interleukin 4 receptor RNA through RNA interference.

Claim 33; SEQ ID NO 389; 128pp; English.

acids (siNAs) which downregulate expression of receptors for interleukingers (e.g., IL-4 receptor (IL-4R), IL-13 receptor (IL-13R) and IL-2 receptor gamma (IL-2RG)) by RNA interference. The invention also relates to similar siNAs which interfere with the expression of the ligands for these receptors, namely IL-4 and IL-13. The siNAs of the invention may or may not comprise ribonucleotides, can contain deoxyribonucleotides, can be chemically modified and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from invention relates to chemically synthesized short interfering nucleic sense oligonucleotide and an antisense oligonucleotide. Specifically,

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the siNAs include short interfering RNA (siRNA), double-stranded RNA, miCro-RNA (miRNA) and short hairpin RNA (shRNA). The invention also cellates to pharmaceutical compositions comprising an siNA targeted to human IL-4R (Refseq accession number NM 000418). IL-13R, IL-2RG, IL-4 or IL-13, especially the siRNAs shown in AEB04245-AEB06055. The invention further discloses expression ovectors and host cells comprising an siNA of the invention. The siNAs exhibit increased resistance to nuclease degradation compared to the prior art. The siNAs of the invention can be used to modulate expression of their target genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of interleukin-related conditions. They may be used in the treatment of cancers and other conditions. They may be used in the treatment of cancers and other conditions, viral infection, inflammatory conditions.

Caucoimmune diseases, respiratory and pulmonary diseases (c.g., asthma, chronic obstructive pulmonary diseases, cand diseases, neurological diseases, neural diseases, neurological diseases, candocrine diseases, neurological diseases, candocrine diseases, prion diseases and cremoning, disenderial diseases, mento diseases, neurological diseases, candocrine diseases, prion diseases and content of diseases, and conditions. The siNAs may also be used in drug screening, diseases, therapeutic target identification, and gene capentic engineering, pharmacogenomics, studying gene function, and gene capenates the sense strand of a human IL-4R-targeted double-stranded continue of a human IL-4R-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  siRNA, which is identical to the human IL-4R transcript target sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 6 A; 8 C; 4 G; 0 T; 1 U; 0 Other;
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Best Local Similarity
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AEB04833 standard; RNA; 19 08-SEP-2005 (first entry) AEB04833; RESULT 1530 AEB04833

ВР.

Human IL-4R siRNA antisense strand, SEQ ID:589.

RNA interference; gene silencing; short interfering RNA; siRNA; cancer; respiratory disease; respiratory gen.; cardiovascular disease; cardiovascular-gen.; neurological disease; neuroprotective; renal disease; endocrine disease; genitourinary disease; nephrotropic; endocrine-gen.; liver disease; genitourinary disease; hepatotropic; ocular disease; ophthalmological; reproductive disease; hepatotropic; antilifertility; gynecology and obstetrics; andrology; mitochondrial disease; prion disease; degeneration; interleukin-4 receptor; IL-4 receptor; ss. hyperproliferation; neoplasm; cytostatic; viral infection; infection; virucide; inflammation; antiinflammatory; autoimmune disease; immunosuppressive; pulmonary disease; 

Homo sapiens

JS2005143333-A1.

30-JUN-2005.

2001US-0306883P. 2001US-0311865P. 2002US-0358580P. 09-JUN-2004; 2004US-00863973 2001US-0292217P 20-JUL-2001; 2 13-AUG-2001; 2 20-FEB-2002; 18-MAY-2001;

2002US-0362016P 2002US-0363124P 2002WO-US015876 06-MAR-2002; 11-MAR-2002; 20-MAY-2002; 셤

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Novel chemically synthesized double stranded short interfering nucleic acid molecule useful for cleaving interleukin 4 receptor RNA through RNA
                                                                                                                                                         Richards I, Polisky B, Mcswiggen J;
                                                                                                                                            (SIRN-) SIRNA THERAPEUTICS INC.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2003US-040129P.
2003US-US00456.
2003WO-US00528.
2003WO-US005346.
2003US-00427160.
                                                              2003US-00444853.
2003US-00693059.
2003US-00720448.
2003US-00727780.
                                                                                          14-JAN-2004; 2004US-00757803.
10-FEB-2004; 2004US-0543480P.
13-FEB-2004; 2004US-00780447.
                                                                                                               16-APR-2004; 2004US-00826966
                                                                                                                             24-MAY-2004; 2004WO-US016390
                                                                                                                                                                       WPI; 2005-457799/46.
                                                              23-MAY-2003; 23-OCT-2003; 24-NOV-2003; 203-DEC-2003;
                            15-JAN-2003;
14-FEB-2003;
                                          20-FEB-2003;
                                                                                                                                                                                                    interference.
                                                         30-APR-2003;
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The invention relates to chemically synthesized short interfering nucleic acids (siNAs) which downregulate expression of receptors for interleukin-

decids (siNAs) which downregulate expression of receptors for interleukin-

decids (lil-ARG) by RNA interference. The invention also relates to similar siNAs which interfere with the expression of the ligands for the heavertion may or the manely IL-ARG) by RNA interfered with the expression of the ligands for the chemically modified and may be double or single stranded. They further comprise sense manely IL-4 and IL-13. The siNAs which minkly and short hairpin RNA (shRNA), double-stranded from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs minkNa) and short hairpin RNA (shRNA). The invention also calters to pharmaceutical compositions comprising an siNA targeted to human IL-4R (RefSeq accession number NM 000418). IL-13R, IL-2RG, IL-4 or IL-13, especially the siNAs shown in AEB04245-AEB06055. The invention of the siNAs exhibit increased resistance to nuclease captants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of interleukin-related conditions. They may be used in the treatment of cancers and other captangen size and conditions. They may be used in the treatment of cancers and other conditions. They may be used in the treatment of cancers and other captangen diseases, respiratory and diseases, prion diseases and creaming diseases, mitochondrial diseases, rend diseases, prion diseases and creaming diseases, mitochondrial diseases, rend diseases, prion diseases and carefully diseases, mitochondrial diseases, rend diseases, prion diseases and carefully diseases, mitochondrial diseases, rend diseases, prion diseases and carefully diseases, mitochondrial diseases, rend diseases, prion diseases and carefully diseases, mitochondrial diseases, rend diseases, prion diseases, prion diseases, production-related conditions. The sinAs may also be used in durance and presents expredu
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Claim 33; SEQ ID NO 589; 128pp; English
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New chemically synthesized double stranded siNA molecule that directs cleavage of an Interleukin-13 receptor (IL-13R) RNA via RNA interference, useful in preparing a composition for treating e.g., inflammatory disorders.
                                                                                                                                                                                                                                          interference; gene silencing; short interfering RNA; siRNA; cancer;
                                                                                                                                                                                                                                                              hyperproliferation; when the state of the st
                                                                                                                                                                                            Human IL-4R siRNA antisense strand, SEQ ID:589.
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                                                AEC14670 standard; RNA; 19 BP
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23-MAY-2003; 2003US-00444853.
23-0CT-2003; 2003US-00693059.
24-NOV-2003; 2003US-0076969.
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2002US-0409293P.
2003US-0440129P.
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10-FEB-2004; 2004US-0543480P.
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2002US-0362016P.
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2004WO-US016390.
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                                                                                                                                                 20-OCT-2005 (first entry)
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17-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-AUG-2005.
                                                                                                AEC14670;
RESULT 1531
                              AEC14670
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/ Match 0.6%; Score 15.8; DB 1; Length 19; Local Similarity 57.9%; Pred. No. 1.2e+03; nes 11; Conservative 6; Mismatches 2; Indels

Query Match

Best Loca Matches

ö

Gaps ö US2005182007-A1.

18-AUG-2005

Homo sapiens.

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Examination of the problem of the properties of the properties of the invention relates to chemically synthesized short interferring nucleic acids (sinNa) which downreallate expression of receptors for interleuking acids (sinNa) which downreallate expression of receptors for interleuking (1.9.7). In receptor (IL-187) by RNA interference. The invention also relates to similar siNNa which interfere with the expression of the ligands for these receptors, namely IL-13 and IL-2. The siNNa of the invention may or may not comprise informed may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, miscontained short interfering RNA (siRNA). double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The invention also relates to pharmaceutical compositions comprising an siNA targeted to human IL-13R (e.g., IL-13R alpha I (IL13RAI), see RefSeq accession number (and pharmaceutical compositions comprising an siNA stayled to the invention can be used to modulate expression of their increased resistance to nuclease degradation compered to the prior art. The siNAs of the invention can be used to modulate expression of their transplants for the transplants for the treatment of a variety of interleukin-related conditions. They may be used in the city of succession of their canders and other proliferative conditions, viral infection, inflammatory conditions, autochmune diseases, respiratory and pulmonary diseases in cells, and transplants for the treatment of a variety of interleukin-related conditions. They may be used in daya gene therapy) and pulmonary diseases, prior diseases, prior diseases, miccolondial diseases, read and expression and validation, genetic employing the signal and expected double-stranded sinna and reproduction-relat
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Query Match

0.6%; Score 15.8; DB 1; Length 19;

Best Local Similarity 57.9%; Pred. No. 1.2e+03;

Matches 11; Conservative 6; Mismatches 2; Indels 0; Gaps

Qy 1280 GGGGCTTTGACATATCT 1298

Db 1 GGGGCUUUGGCAUGUCCU 19

RESULT 1532
AEC14470/C
ID AEC14470 standard; RNA; 19 BP
XX
AC AEC14470;
XX
DT 20-OCT-2005 (first entry)
XX
DY
XX
DW Interference; gene silenc:
XX
XW
Nyperproliferation; neoplasm;
XW
Nyperproliferation; neoplasm;
XW
Nyperproliferation; neoplasm;
XW
Nymune disorder; immunosuppre
XW
remal disease; respirate
XW
renal disease; endocrine disease;
XW
endocrine-gen.; interlogism;
XW
coular disease; ophthalmologism;
XW
coular disease; ophthalmology an
XW
interleukin-4 receptor; IL-4;
XW
interleukin-4 receptor; IL-4

RNA interference; gene silencing; short interfering RNA; siRNA; cancer; hyperproliferation; neoplasm; cytostatic; viral infection; infection; virucide; inflammation; antiinflammatory; autoimmune disease; immunosuppressive; pulmonary disease;

Human IL-4R transcript target sequence/siRNA sense strand, SEQ ID:389.

immune disorder; immunosuppressive; pulmonary disease; respiratory disease; respiratory disease; respiratory-gen.; cardiovascular disease; cardiovascular-gen.; neurological disease; neuroprotective; renal disease; endocrine disease; gentrourinary disease; nephrotropic; endocrine-gen.; liver disease; gastrointestinal disease; hepatotropic; ocular disease; ophthalmological; reproductive disorder; infertility; anneinfertility; gynecology and obsettrics; andrology; mitochondrial disease; prion disease; degeneration; interlibity interleukin-4 receptor; IL-4 receptor; ss.

(SIRN-) SIRNA THERAPEUTICS INC 2001US-0306883P. 2001US-0311865P. 2002US-0358580P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 39-JUN-2004; 2004US-00863973. 2002US-0362016P. 2002US-0363124P. 14-FEB-2003; 2003WO-US004566. 20-FEB-2003; 2003WO-US005028. 20-FEB-2003; 2003WO-US005346. 20-AUG-2004; 2004US-00922675 2002US-0386782P 2003US-0440129P 2003US-00720448 14-JAN-2004; 2004US-00757803. 10-FEB-2004; 2004US-0543480P. 2003US-00427160 03-DEC-2003; 2003US-00727780 2003US-00444853 Mcswiggen J, Beigelman L; 4PI; 2005-581759/59 29-AUG-2002; 2 05-SEP-2002; 2 09-SEP-2002; 2 15-JAN-2003; 30-APR-2003; 06-MAR-2002; 20-FEB-2002; 06-JUN-2002; 23-MAY-2003; 24-NOV-2003; 24-MAY-2004;

New chemically synthesized double stranded siNA molecule that directs cleavage of an interleukin-13 receptor (IL-13R) RNA via RNA interference, useful in preparing a composition for treating e.g., inflammatory disorders.

Claim 33; SEQ ID NO 389; 127pp; English.

The invention relates to chemically synthesized short interfering nucleic caids (siNAs) which doursegulate expression of receptors for interleukin-caids (siNAs) which doursegulate expression of receptors for interleukin-lessing and in-3 receptor (IL-4R) and IL-2 receptor gamma (IL-2RG)) by RNA interference. The invention also relates to similar siNAs which interfere with the expression of the ligands for these receptors, namely IL-13 and IL-4. The siNAs of the invention may or these receptors, namely IL-13 and IL-4. The siNAs of the invention may or comprise ribonucleotides, can contain deoxyribonucleotides, can comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, can ense oligonucleotide short hairpin RNA (siRANA), double-stranded RNA, cancer kopharmaceutical compositions comprising an siNA (shanA). The invention also calls comprising an siNA of the invention rule siNAs shown in concased resistance to nuclease degradation compared to the prior art. The siNAs of the invention can be used to modulate expression of their carget genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a cancers and other proliferative conditions, viral infection,

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diseases (e.g., asthma, chronic obstructive pulmonary disease (COPD), allergies), cardiovascular diseases, neurological diseases (COPD), diseases, cardiovascular diseases, neurological diseases, renal diseases, ocular diseases, liver diseases, mitochondrial diseases, endocrine diseases, prion diseases and reproduction-related conditions. The slMAs may also be used in drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the sense strand of a human IL-4R-targeted double-stranded siRNA, which is identical to the human IL-4R transcript target sequence. Note: The sequence data for this patent is also available in electronic format directly from the US patent of the contract of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NOGO receptor; RNA interference; gene silencing; neuroprotective; gene therapy; neurological disease; short interfering RNA; siRNA; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NOGO receptor target region/siRNA molecule sense strand, SEQ ID 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                            segdata.uspto.gov/sequence.html?DocID=20050182007.
                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 6 A; 8 C; 4 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1280 GGGGCTTTGACATATCCT 1298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 GGGGCTTTGGCATGTCCT 1
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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2002US-0358580P.
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2002WO-US015876.
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2003US-00727780.
2004US-00757803.
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Best Local Similarity 89.5
Matches 17; Conservative
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11-MAR-2002;
03-APR-2002;
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18-MAY-2001;
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20-FEB-2002;
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                                                                                                                                                                                                      New chemically synthesized double stranded siNA molecule that directs cleavage of a NOGO receptor RNA via RNA interference, useful in preparing a composition for treating e.g., neurological disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NOGO receptor; RNA interference; gene silencing; neuroprotective; gene therapy; neurological disease; short interfering RNA; siRNA; ss.
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Pred. No. 1.2e+03;
2; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 14 A; 1 C; 0 G; 0 T; 4 U; 0 Other;
                                                                                                                                    Chowrira BM, Haeberli P;
                                                                                                                                                                                                                                                                        Claim 33; SEQ ID NO 99; 192pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2705 TACTAAAAAAAAAAAA 2723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   siRNA molecule of the invention.
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09-FEB-2001; 2001US-00804273.

05-ARR-2001; 2001US-00827395.

18-MAY-2001; 2001US-0306883P.

13-AUG-2001; 2001US-0368889P.

20-FEB-2002; 2002US-0358880P.
                                                                                                   (SIRN-) SIRNA THERAPEUTICS INC.
              13-FEB-2004; 2004US-00780447.
16-APR-2004; 2004US-00826966.
30-APR-2004; 2004WO-US013456.
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                                                                2004WO-US016390
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Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                     WPI; 2005-581760/59.
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                                                                                                                                     Mcswiggen J,
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                                                                  24-MAY-2004;
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Claim 33; SEQ ID NO 99; 206pp; English.
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                                                                                                                                                                24-NOV-2003; 2003US-00720448.
03-DEC-2003; 2003US-00727780.
                                                                                                                                                                                       14-JAN-2004; 2004US-00757803.
10-FEB-2004; 2004US-0543480P.
13-FEB-2004; 2004US-00780447.
16-APR-2004; 2004US-00826966.
                                                                                                                                                                                                                                                  24-MAY-2004; 2004WO-US016390.
                                                                                                                              20-AUG-2004; 2004WO-US026930
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                                                                                                                                                                                                                                                                                                                       WPI; 2005-746892/76.
                                                                                                                                                                                                                                                                                                                                                                                 disease or dementia.
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                                                                                  WO2005045035-A2
                                                                                                                                                     23-OCT-2003;
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                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel chemically synthesized double stranded short interfering nucleic acid (siNA) molecule which directs cleavage of a NOGO receptor RNA via RNA interference (RNAi). Each strand of the siNA molecule is 18-23 nucleotides in length. One strand of the siNA molecule NOGO receptor RNA. The siNA molecules include short interfering RNA (siRNA), double-stranded RNA (dsRNA), micro-RNA (miRNA) and short hairpin molecules (siRNA). The invention also includes a composition comprising the siNA molecule in a carrier or diluent. The siNA molecules have neuroprotective activity and may be useful in gene therapy. The double stranded siNA molecules are useful in preparing a composition for treating NOGO receptor-associated disorders, e.g. neurological disorders. This sequence represents a NOGO receptor siRNA molecule antisense strand
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                                                                                                                                                                                                                                                                                                                                                      New chemically synthesized double stranded siNA molecule that directs cleavage of a NOGO receptor RNA via RNA interference, useful in preparing a composition for treating e.g., neurological disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.6%; Score 15.8; DB 1; Length 19; 89.5%; Pred. No. 1.2e+03;
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0; Mismatches 2
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                                                                                                                                                                                                                                                                                                         Chowrira BM, Haeberli P;
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                                           2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                     2002WO-US010512.
2002WO-US015876.
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2003WO-US005346.
                                                                                                                             2003US-00427160.
2003US-00430882.
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                                                                                            2003US-0440129P.
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2003US-00693059.
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2004WO-US013456
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Best Local Similarity 89.5'
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                                                                                          15-JAN-2003;
20-FEB-2003;
20-FEB-2003;
                                           06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                            30-APR-2003;
06-MAY-2003;
23-MAY-2003;
23-OCT-2003;
                                                                                                                                                                                                                                                                                                         Mcswiggen J,
                     03-APR-2002;
17-MAY-2002;
                                                                                09-SEP-2002;
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03-DEC-2003;
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The invention describes a chemically synthesized double-stranded short interfering nucleic acid (siNA) molecule (I) directing cleavage of NOGO creceptor RNA through RNA interference (RNAi), where each strand of the molecule is 18-23 nucleotides in length, and one strand of the molecule comprises nucleotide sequence having sufficient complementarity to the NOGO receptor RNA for the siNA molecule to direct cleavage of the NOGO receptor RNA through RNAi. Also described is a composition comprising (I) together with a carrier or diluent. (I) is useful for downregulation or inhibition of expression of NOGO and/or NOGO receptor haplotype polymorphism that are associated with a disease or condition (e.g., neurologic diseases, disorders and/or conditions). (I) is useful for treating or preventing central nervous system (CNS) injury, cerebrovascular accident (CVA), stroke, Alzheimer's disease, dementia or multiple sclerosis. (I) is useful as reagents in ex vivo applications e.g., in tissue or cells that are transplanted into a subject for therapeutic effect. This sequence represents a NOGO receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of NOGO (Neurite outgrowth inhibitor) receptor RNA through RNA interference, useful in treating Alzheimer's
antisense therapy; gene expression; neurological disease; nervous system injury; cerebrovascular ischemia; Alzheimers disease; dementia; multiple sclerosis; NGO receptor; siRNA; short interfering RNA; RNA interference; gene silencing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 14 A; 1 C; 0 G; 0 T; 4 U; 0 Other;
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The invention relates to a method of identifying a compound that induces chondrocyte anabolic stimulation. The methods and agent are useful for treating and/or preventing a disease involving a systemic or local carease in cartilage, e.g. osteoarthritis, rheumatoid arthritis, septimic or infectious arthritis, reactive arthritis, gouty arthritis, septimic or infectious arthritis, reactive arthritis, reflex sympathetic septic or infectious arthritis, reactive arthritis, reflex sympathetic intropathy, algodystrophy, Tietze syndrome or costal chondritis, intropathy, osteoarthritis deformans endemica, Meeleni disease, Handigodu disease, degeneration resulting from fibromyalgia, systemic lupus erythematosus, scleroderma, ankylosing spondylitis, hereditary chondrolysis, chondrodysplasia, pseudoachondrodysplasias, microtia, and metaphyseal chondrodysplasia. The agent is useful in the manufacture of a medicament for the treating and/or preventing a disease involving a decrease in mean cartilage thickness, e.g. osteoarthritis, chypercalcemia of malignancy, multiple myelomatosis, hyperparathyroidism, and hyperthyroidism. The present sequence represents a human chondrocyte anabolic stimulation associated target DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a compound that induces chondrocyte anabolic stimulation, useful for treating osteoarthritis, comprises measuring a compound-polypeptide property related to the anabolic stimulation of chondrocytes.
                                                                                                                  Osteopathic; Nootropic; Neuroprotective; Dermatological;
Antiinflammatory; Antiarthritic; Antiarthritic; musculoskeletal disease;
chondrocyte anabolic stimulator; ss.
                                                                   Human chondrocyte anabolic stimulation target DNA SEQ ID NO 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.6%; Score 15.8; DB 1; Length 19; 89.5%; Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tomme PHM, Michiels F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 4; SEQ ID NO 352; 179pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTGAAGAAGCTGGAGTTGC 19
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                                                                                                                                                                                                                                                                                                                                                                               21-JUN-2005; 2005WO-EP052875.
                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUN-2004; 2004US-0581568P.
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                       09-MAR-2006 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GALA-) GALAPAGOS NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2006-067565/07.
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                                                                                                                                                                                                                                                                            WO2005124342-A2.
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                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                           29-DEC-2005.
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV19118,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a chemically synthesized double-stranded short interfering nucleic acid (siNA) molecule (I) directing cleavage of NOGO receptor RNA intrough RNA interference (RNAi), where each strand of the molecule is 18-23 nucleotides in length, and one strand of the molecule comprises nucleotide sequence having sufficient complementarity to the NOGO receptor RNA for the siNA molecule to direct cleavage of the NOGO receptor RNA through RNAi. Also described is a composition comprising (I) together with a carrier or diluent. (I) is useful for downregulation or inhibition of expression of NOGO and/or NOGO receptor proteins arising (from NOGO and/or NOGO receptor proteins arising in a disease or conditions). (I) is useful for treating or preventing central nervous system (CNS) injury, cerebrovascular accident (CNA), stroke, Alzhaimer's gisease, dementia or multiple sclerosis. (I) is useful as reagents in exvivo applications e.g., in tissue or cells that are transplanted into a subject for therapeutic effect. This sequence represents a NOGO receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel chemically synthesized double-stranded short interfering nucleic acid molecule directing cleavage of NOGO (Neurite outgrowth inhibitor) receptor RNA through RNA interference, useful in treating Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                      antisense therapy, gene expression, neurological disease, nervous system injury; cerebrovascular ischemia; Alzheimers disease, dementia, multiple sclerosis; NOGO receptor; siRNA; short interfering RNA; RNA interference; gene silencing; ss.
                                                      CNS-gen.; neuroprotective; nootropic; cerebroprotective; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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89.5%; Pred. No. 1.2e+03;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 4 A; 0 C; 1 G; 0 T; 14 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 33; SEQ ID NO 198; 206pp; English.
Human NOGO receptor siRNA SEQ ID NO 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2705 TACTAAAAAAAAAAAA 2723
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2003US-00727780.
2004US-05757803.
2004US-0573480P.
2004US-00780447.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEF14493 standard; DNA; 19 BP
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                                                                                                                                                                                                                                                                                                                                                        20-AUG-2004; 2004WO-US026930
                                                                                                                                                                                                                                                                                                                                                                                                            2003US-00693059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2004WO-US013456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen J, Chowrira BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 89.5
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease or dementia
                                                                                                                                                                                                                                                       WO2005045035-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JAN-2004;
10-FEB-2004;
                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                            23-OCT-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                      24-NOV-2003;
03-DEC-2003;
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Mille-Baker B;

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Indels

Query Match

δ 엄 AEF14493

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AEF14493

Secreted apoptosis-related protein; SARP; mSARP1; mouse; prostate cancer;

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WPI; 2000-257789/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-APR-1999;
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23-JUN-1998;
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                                                                                                                                                                                                                                                                                                            AAA25453;
                                                                                                                                                                                                                                                                       RESULT 1540
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Matches
                                                                                                                                                                                                                                                                               AAA25453,
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                                                                                                                                                                                                                                     This oligo(T) synthetic oligonucleotide was used for first strand cDNA synthesis from total RNA isolated from either logarithmically growing or quiescent 10T1/2 mouse fibroblast cells. It was also used with an arbitarary d(N10) primer in PCR. The PCR products were used in a differential display to identify the mSARP1 gene (see AAV19112) that codes for novel murine secreted apposis-related protein mSARP1 (see AAW3914). The invention relates to SARP polynucleotides (see also AAW19113-15) and polypeptides (see also AAW19113-15), antibodies specific and therapeutic methods, and methods for treating diseases related to the regulation of SARP expression in tissue and body fluid samples, including
                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                               New secreted apoptosis-related proteins - useful for modulating apoptosis, particularly for treatment of prostatic or breast cancer, also for diagnosis and monitoring of disease.
                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 15.6; DB 1; Length 17; 88.2%; Pred. No. 1.2e+03; ive 1; Mismatches 1; Indels
breast cancer; diagnosis; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amplification; detection; gene expression; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HMRI ) HOECHST MARION ROUSSEL DEUT GMBH.
                                                                                                                                                                                                                      Example 1; Page 30; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                          2707 CTAAAAAAAAAAAA 2723
                                                                                                                         (LXRB-) LXR BIOTECHNOLOGY INC.
                                                                                            96US-0026603P.
96US-0028363P.
                                                                          97WO-US017154
                                                                                                                                                                                                                                                                                                                                                                                                                                    98DE-01040731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ89372 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                            Melkonyan H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA detecting primer #2
                                                                                                                                                             WPI; 1998-230704/20.
                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 15; Conserv
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                                                                                            24-SEP-1996;
11-OCT-1996;
                                   WO9813493-A2
                                                                         24-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
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                                                                                                                                            Umansky S,
                  Synthetic
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                                                                                                                                                                                                          which comprises amplifying cDNA with first and second differently labeled primers and analysis of the amplified labeled cDNA. The method is useful for analyzing differential gene expression, for identifying and/or characterizing pharmacological activities or for identifying target genes. The use of different primer combinations allow more cDNAs to be amplified. The method also provides a more detailed analysis than prior art methods. This sequence represents a primer used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes nucleic acids (A) that interact stably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1951.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonuclectide; gene expression modification; cancer; bosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
Analysis of RNA samples, useful for detection of differential gene expression uses two differently labeled primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bellon L;
                                                                                                                                                                            invention describes a novel method for analysis of an RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.6; DB 1; Length 17;
Pred. No. 1.2e+03;
1; Mismatches 0; Indels
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Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Beigelman L, Mcswiggen JA,
Zwick M, Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids that interact, and used to treat cancer.
                                                                                                   Disclosure; Page 10; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 77; Page 79; 148pp; English.
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98US-00103636.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA25453 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             method of the invention
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Best Local Similarity
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                         can also be used to
for other conditions associated with levels of oestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent coetrogen receptor hammerhead riboxyme sequences. AAA2593 to AAA2593 to AAA2593 to estream their corresponding target sequences.
                                                                                                                                                                                             sequences, and AAA26107 to AAA26218 represent their corresponding target sequences. AAA26219 to AAA26271 represent other ribozyme sequences and antisense oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acids that interact, and optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention describes nucleic acids (A) that interact stably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                / Match 0.6%; Score 15.4; DB 1; Length 17; Local Similarity 94.1%; Pred. No. 1.28+03; Hes 16; Conservative 0; Mismatches 1. *-1.1.
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Haeberli P;
                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
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Zwick M, Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 77; Page 79; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                             2705 TACTAAAAAAAAAAA 2721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA25452 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to treat cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matulic-Adamic J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-APR-1999;
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23-JUN-1998;
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                                                                                                                                                                                                                                                            invention
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Best Local S:
Matches 16
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correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research usegenes (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves AAA2447 represent oserrogen receptor hammerhead ribozyme sequences, and AAA2447 represent oserrogen receptor hammerhead ribozyme sequences, and AAA2593 represent their corresponding target sequences. AAA2593 to AAA26105 represent their corresponding target sequences, and AAA26107 to AAA2611 represent their corresponding target sequences. Sequences. AAA26219 to AAAA2671 represent their corresponding target sequences and antisense oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probes for detecting target nucleotide sequence in sample, has sequence that forms hairpin structure having a double-stranded segment and singlestranded loop collectively forming region complementary to target
                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                             Length 17;
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                                                                                                                                                                                                                                        BP; 0 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                         0.6%; Score 15.4; DB 1;
14.1%; Pred. No. 1.2e+03;
.ve 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENE-) APPLIED GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cocation/Qualifiers
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30-MAR-2001; 2001US-00823647.
                                                                                                                                                                                                                                                                                            94.1%;
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/label= RNA
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                                                                                                                                                                                                                                                                                                                16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-2002
                                                                                                                                                                                                                                          Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                          invention
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ABA91530/
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number for RNA cleavage. The invention provides probes for nucleic acid hybridisation. The probes form a hairpin structure comprising a double-stranded stem and a single-stranded loop, and are capable of both intramolecular and intermolecular hybridisation. The double-stranded stem may comprise a methylphosphonate DNA:RNA hybrid that is resistant to strand in the DNA:RNA duplex becomes sensitive to RNAse H treatment and strand in the DNA:RNA duplex becomes sensitive to RNAse H treatment and can be removed. Arrays and methods for nucleic acid hybridisation using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method of sequential consensus region-directed amplification for sorting a mixture of DNAs into 2 or more subsets or distinguishing gene expression patterns in 2 samples. The methods, kits and oligonucleotides are useful for sorting a mixture of DNAs into 2 or more subsets or distinguishing gene expression patterns in 2 samples e.g. for disease diagnosis and gene analysis. The present sequence is oligo APPCR primer used to illustrate the method of the invention
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in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequential consensus region-directed amplification; gene expression; disease diagnosis; gene analysis; human; matrix metalloproteinase; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequential consensus region-directed amplification for sorting mixture DNAs into 2 or more subsets or distinguishing gene expression patterns 2 samples, useful for disease diagnosis and gene analysis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.6%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03; iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 1 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD44151 standard; DNA; 17 BP.
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97US-0108152P,
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                                                                                                                                                                                                                                                                                                           the probes are provided
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Matches 16; Conserv
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p2.0.2. and MD212 is encoded at chromosome 16p11.2 and MD212 is encoded at chromosome 15p2.0.2. and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23. MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                              Cytostatic, immunostimulant, gene therapy, vaccine, human, zinc finger protein; MDZ3; MDZ4, MDZ12; chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.6%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03; ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                            Human MDZ7 scanning oligonucleotide SEQ ID 5255.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 5255; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2709 AAAAAAAAAAAAA 2725
                                                                                            ADB04269 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB04273/c
ID ADB04273 standard; DNA; 17 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gu Y, Nguyen C;
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17 AAAAAAAAAAAABA
                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                           developmental disorder; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AEOM-) AEOMICA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-423107/40
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                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                ADB04269;
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Query Match
Best Local Similarity 94.1
Matches 16; Conservative

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Gaps

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1; Indels

0; Mismatches

0.6%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is conceded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 16p1) 2 and MD212 is encoded at chromosome 16p1) 2 and MD212 is encoded at chromosome 16p1) 2 and MD212 is encoded at chromosome 16p2. MD27, and MD212 is encoded at chromosome 16p2 in an anufacturing a medicament for treating or preventing a disorder as associated with decreased or increased expression or activity of MD23. MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins. The present sequence was used to illustrate the invention.
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                                                                   Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; Chromosome 7q22.1; chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ12; chromosome 7q22.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.6%; Score 15.4; DB 1; Length 17;
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                                                   Human MDZ7 scanning oligonucleotide SEQ ID 5259.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; SEQ ID NO 5259; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                           MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                 02-AUG-2001; 2001US-00922181
                                                                                                                                                                                                                                                                                                    Shannon M, Gu Y, Nguyen C;
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Best Local Similarity 94.1%;
Matches 16; Conservative
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                          (first entry)
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                                                                                                                                              Homo sapiens
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                            20-NOV-2003
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  ADB04273
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is concoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p1.2 and MDZ12 is encoded at chromosome 16p2.2 and MDZ12 is encoded at chromosome 16p2.2 and MDZ12 is encoded at chromosome 16p2.2 and MDZ12 is encoded. The nucleic associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids and also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes for detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapoutic agents for gene therapy or as proteins are useful as therapoutic agents for gene therapy or as
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chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; SEQ ID NO 5260; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gu Y, Nguyen C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AEOM-) AEOMICA INC.
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                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB04270;
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is encoded at chromosome 7q22.1, MD24 is encoded at chromosome 6p21.3-22.2, MD27 is encoded at chromosome 6p21.3-22.2, MD24, MD24, MD24, MD27, and MD212 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders: The nucleic caused by altered expression of MD23, MD24, MD27, or MD212, e.g. cancer of detect and characterize gross active and also be used as probes to detect and characterize gross alterations in MD23, MD24, MD27, or MD212 genetic locus. The probes are useful in constructing microaarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                 New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.6%; Score 15.4; DB 1; Length 17; Best Local Similarity 94.1%; Pred. No. 1.2e+03; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17: BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                     Example 8; SEQ ID NO 5256; 103pp; English.
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                    30-JUL-2002; 2002EP-00016874.
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                                                           02-AUG-2001; 2001US-00922181.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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ABZ61225 standard; RNA; 17
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                                                                                                                                         Nguyen
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                                                                                                  (AEOM-) AEOMICA INC.
                                                                                                                                         Gu Y,
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                                                                                                                                       Shannon M,
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing ALSO useful for treating breat, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and ALDS. The sequences shown in ABZ59889 - ABZ65216, ABZ65511, ABZ6550 - ABZ66524, ABZ66530 - ABZ6658 represent substrate/target sequences for the human ribozymes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CS injury; epinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupue; multiple sclerosis; transplant rejection;
                                                                Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     0.6%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 2 A; 4 C; 8 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.1%; Pred. .v.
tive 0; Mismatches
                                                                                                                                     Claim 58; Page 111; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PKR substrate sequence #522.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1737 ACCTGAGGCCCTCCCTG 1753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCAGAGGCCCTCCCTG 1
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 94.1
nes 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200281628-A2
 Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 substrate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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WPI; 2003-058513/05

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Gaps

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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NoGO, prostaglandin D2 receptor (PTGDR),
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
confined or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischaemia/reperfusion injury, glomerulomephritis, sepsis, and allergic
conditions (e.g. asthma, allergic thinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifte and mutations within diseased cells or to detect the presence of
target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide; neurite growth.inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; prostal kinase PKR; cerebrovascular accident; central kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoride accidential autoimmune disease; lupus; multiple solerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PKR;
Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 0.6%; Score 15.4; DB 1; Length 17; Local Similarity 94.1%; Pred. No. 1.2e+03; les 16; Conservative 0; Mismatches 1; Indels
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                                                                                  Claim 59; SEQ ID NO 2941; 317pp; English
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       substrate sequence.
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Matches
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that down regulate the expression or inhibit the function of a receptor for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PrGDR), ISAPPAB kinase (IKK), or protein kinase PKR. The nucleic acids of the invention are useful for treating: cerebrowascular accident, central nervous system (CMS) injury, spinal cord injury, cancer (e.g. melanoma, CT lymphoma or glioma), inflammatory disease (e.g. relamaciól arthritis, restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The nucleic acids of the invention are also useful for down-regulating the cypression of a target gene and as a diagnostic tool to examine genetic diffs and mutations within diseased calls or to detect the presence of a target con an acell. The present RNA sequence represents a human PKR ö CpG immunostimulatory oligonuclectide; immune response; allergy; asthma; viral infection; bacterial infection; cancer; lymphoma; intraepithelial neoplasm; melanoma; neuroblastoma; Hodgkin's lymphoma; carcinoma; sarcoma; gene therapy; phosphorothioate; ss. Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease. Gaps .; 0 Length 17; 1; Indels Sequence 17 BP; 0 A; 1 C; 0 G; 0 T; 16 U; 0 Other; /note= "Phosphorothioate backbone" 0.6%; Score 15.4; DB 1; 94.1%; Pred. No. 1.2e+03; ative 0; Mismatches 1; Uhlmann E; CpG immunostimulatory oligonucleotide #11. SEQ ID NO 2940; 317pp; English. Location/Qualifiers /\*tag= a /mod\_base= OTHER 2709 AAAAAAAAAAAAAA 2725 Krieg AM, Jurk M, Vollmer J, (COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH. ADP86140 standard; DNA; 17 BP 11-DEC-2003; 2003WO-US039775. 11-DEC-2002; 2002US-0432409P. 25-SEP-2003; 2003US-0506108P. 17 AAAAAAAAAAAAAGAA 1 09-SEP-2004 (first entry) Conservative WPI; 2004-487902/46 substrate sequence. Local Similarity WO2004053104-A2 Key modified base Unidentified 24-JUN-2004. 16; ADP86140; Query Match RESULT 1551 Matches ADP86140/ ઠે 셤 

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New oligonucleotides, useful for treating allergy or asthma, viral and
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                                                                             Query Match
Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                   Key
modified_base
                                                                                                                                                             Unidentified
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                                                                                                                         ADP86146;
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Query Match
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                                                                                                                                           The invention relates to a class of CpG immunostimulatory
oligomucleotides containing a 5'TCG motif or a CG at or the 5' end that
are useful for stimulating an immune response. Oligomucleotides and
compositions of the invention are useful for treating allergy or asthma,
viral and bacterial infections and cancer e.g. biliary tract cancer,
breast cancer, cervical cancer, choriocarcinoma, colon cancer,
endometrial cancer, gastric cancer, lymphomas, intraepithelial neoplasms,
cerdometrial cancer, gastric cancer, pancreatic cancer, prostate cancer,
certal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer,
cetal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer,
and CNS cancer, connective tissue cancer, osesophageal cancer, eye cancer,
cetal cancer, as well as other cavity cancer, skin cancer,
testicular cancer, as well as other carcinomas and sarcomas. The
invention is also useful in gene therapy. The present sequence is a CpG
immunostimulatory oligonucleotide.
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New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other;
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                                                                                                      Example; SEQ ID NO 11; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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ADP86146 standard; DNA; 17 BP.
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25-SEP-2003; 2003US-0506108P.
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                                                    cancer, cervical cancer.
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The invention relates to a class of CpG immunostimulatory oligonucleotides containing a 5'TCG motif or a CG at or the 5' end that are useful for stimulating an immune response. Oligonucleotides and compositions of the invention are useful for treating allersy or asthma, viral and bacterial infections and cancer e.g. biliary tract cancer, breast cancer, gastric ancer, thoriocarcinoma, colon cancer, endometrial cancer, gastric ancer, lymphomas, intraspithelial neoplasms, liver cancer, lung cancer (e.g. small cell and non-small cell), melanoma, neuroblastomas, ovarian cancer, pancreatic cancer, prostate cancer, rectal cancer, sarcomas, thyroid cancer, renal cancer, bone cancer, brain and CNS cancer, connective tissue cancer, rosophageal cancer, brain and CNS cancer, as well as other carcinomas and sarcomas. The invention is also useful in gene therapy. The present sequence is a CpG incorrection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
bacterial infections, and cancer, e.g. biliary tract cancer, breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 0.6%; Score 15.4; DB 1; Length 17; Local Similarity 94.1%; Pred. No. 1.2e+03;
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/note= "Phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 0 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
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                                                         ID NO 17; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                            immunostimulatory oligonucleotide.
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25-SEP-2003; 2003US-0506108P.
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/label= RNA
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                   cancer, cervical cancer.
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modified_base
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WPI; 2004-487902/46

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oligonucleotides containing a S'TCG motif or a CG at or the S' end that
are useful for stimulating an immune response. Oligonucleotides and
compositions of the invention are useful for treating allergy or asthma,
viral and bacterial infections and cancer e.g. biliary tract cancer,
breast cancer, cervical cancer, choriocarcinoma, colon cancer,
endometrial cancer, gastric cancer, lymphomas, intraepithelial neoplasms,
liver cancer, lung cancer (e.g. small cell and non-small cell), melanoma,
neuroblastomas, ovarian cancer, pancreatic cancer, prostate cancer,
rectal cancer, sarcomas, thyroid cancer, renal cancer, brain
and CNS cancer, connective tissue cancer, respandable cancer, brain
and CNS cancer, as recomes, thyroid cancer, respandable cancer, brain
testicular cancer, as well as other carcinomas and sarcomas. The
invention is also useful in gene therapy, The present sequence is a CpG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CpG immunostimulatory oligonucleotide; immune response; allergy; asthma; viral infection; bacterial infection; cancer; lymphoma; intraeptthelial neoplasm; melanoma; neuroblastoma; Hodgkin's lymphoma; carcinoma; sarcoma; gene therapy; phosphorothioate; DNA-RNA hybrid; ss.
    New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast cancer, cervical cancer.
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                                                                                                                                                                                                                                                                                                Match 0.6%; Score 15.4; DB 1; Length 17; Local Similarity 94.1%; Pred. No. 1.2e+03; les 16; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                 immunostimulatory oligonucleotide (DNA-RNA hrbrid)
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                                                         Example, SEQ ID NO 56, 104pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP86184 standard; DNA; 17 BP
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25-SEP-2003; 2003US-0506108P.
                                                                                                                                                                                                                                                                                                                                                                        17 AAAAAAAAAAAAAAA 1
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/label= RNA
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/*tag= a
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP86184;
                                                                                                                                                                                                                                                                                                   Query Match
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Matches
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                                                                                   New oligonucleotides, useful for treating allergy or asthma, viral and bacterial infections, and cancer, e.g. biliary tract cancer, breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         short interfering RNA; siRNA; RNA interference; gene silencing; cytostatic; cancer; Ras gene; substrate; ss.
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                                                                                                                                                                                                                                                                                                                                                            The invention relates to a class of CpG immunostimulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 17 BP; 0 A; 1 C; 0 G; 15 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human H-Ras substrate RNA sequence SEQ ID NO:1337.
                                                                                                                                                                                                                                                                        Example; SEQ ID NO 55; 104pp; English.
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2001US-0296249P.
2001US-0306883P.
2001US-0311865P.
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2002US-00157580.
2002WO-US016840.
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2002US-0358580P.
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2002US-0363124P.
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2002US-0386782P.
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                                                                                                                                                                           cancer, cervical cancer.
WPI; 2004-487902/46.
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05-SEP-2002;
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20-FEB-2002;
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The invention relates to a double-stranded short interfering RNA (siRNA)

molecule (1) comprising a first nucleotide sequence having 19-23

nucleotides complementary to an RNA sequence encoding HRR2 or its

portion, and a second nucleotide sequence, having 19-23 nucleotides

contion, and a second nucleotide sequence, and including at least

one nucleotide that is not a 2'-OH containing ribonucleotide. Also

described is a method of producing a class of nucleic acid-based gene

modulating agents that exhibit a high degree of specificity for RNA of a

desirbed target. (I) is useful for modulating HRR2 activity in a cell, and

containing agents that exhibit a high degree of specificity for RNA of

desirbed target. (I) is useful for modulating the RR2 activity in a cell, and

containing acancer, breast cancer, such as pancreatic cancer,

bladder cancer, lung cancer, breast cancer or prostrate cancer. The

present sequence represents a human H-Ras substrate RNA sequence for a

NNAzyme (ribozyme), which is used in an example from the present

invention for the identification of potential target sites in human Ras
                                                                                                                                                                                                                                                                                                                                                                                  Novel double-stranded short interfering RNA molecule having first nucleotide sequence complementary to RNA encoding HER2 or its portion, and second nucleotide sequence having complementarity to first sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; SEQ ID NO 1337; 143pp; English.
                                                                                                                                                                                                                                                          (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                     16-APR-2003; 2003US-00417012.
24-APR-2003; 2003US-0042704.
30-APR-2003; 2003US-00427160.
23-MAY-2003; 2003US-00444883.
                                                              2003WO-US005028.
                                                                                                                                                                                                                      2003US-00693059.
  2002US-0409293P
                                            2003US-0440129P
                                                                                                                                                                                              2003US-00652791
                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for treating cancer.
                                                                                                                                                                                                                                                                                                                                              WPI; 2005-331166/34.
                                                                                                                                                                                                                  23-OCT-2003;
                                                                                    20-FEB-2003;
                                                                                                                                                                                              29-AUG-2003;
                                                                20-FEB-2003;
                                                                                                                                                                                                                                                                                                    Mcswiggen J;
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Ouery Match 0.6%; Score 15.4; DB 1; Length 17; Best Local Similarity 94.1%; Pred. No. 1.2e+03; Sequence 17 BP; 2 A; 4 C; 8 G; 0 T; 3 U; 0 Other;

1737 ACCTGAGGCCCTCCCTG 1753 17 ACCAGAGGCCCTCCCTG 1 용 ð

AED81301 standard; DNA; 17 (first entry) 26-JAN-2006 AED81301; RESULT 1556 AED81301 

IL-10 expression assay, test oligonucleotide SEQ ID No:59.

pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.

Synthetic.

VO2005111057-A2

24-NOV-2005

04-APR-2005; 2005WO-US011827.

02-APR-2004; 2004US-0558951P.

(COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH. (COLE-)

Krieg AM, Vollmer J;

WPI; 2005-786756/80.

New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus.

Example; SEQ ID NO 59; 111pp; English

The invention relates to an oligonucleotide having the formula: (a) 5' XIXIXZX2 3' where 5' denotes the 5' and of the oligonucleotide and where 5' and cotted the oligonucleotide, where X is a T or modified The concloside, and where a 3' of an olification the concloside and Where a 3' of an olification and Na and Na are polymucleotide that do not include a CG dinucleotide, and Where a 3' of the oligonucleotide and the NZ dinucleotide and where a 3' olification and Na and NZ are polymucleotide and the NZ dinucleotide and where a 3' olification and NZ are polymucleotide and where a 3' olification and NZ are polymucleotide and where a 3' olification of the oligonucleotide in the oligonucleotide and where NZ is a core modified The nucleotide. NI is a composition to nucleotide, and where NZ is a composition to composition the size of the month of the oligonucleotide of 5 to 30 nucleotides, where NI does not include 5' a nucleotide are (1) a pharmaceutical composition for onerginal matibodies, and where NZ is polymucleotide of 5 to 30 nucleotides, and where NZ is polymucleotide of 5 to 30 nucleotides, and office and manaceutical composition comprising the oligonucleotide in combination with a therapeutic agent eslected from chancherapeutic agents encolonal matipoderic and where NZ is pharmaceutical composition to comprising an anticonal matipode or application of supplied to a palative to interferon (ITM) alpha expression in a subject. comprising administering an immunostiminatory nucleic and composition to a subject where the subject in the subject in a subject by administering an immunostiminatory nucleic acid or composition to the subject where the mannostiminatory nucleic acid or composition to the subject where the immunostiminatory nucleic acid or composition to the subject where the immunostiminatory nucleic acid or composition to the subject where the immunostiminatory nucleic acid or composition to the subject where the immunostiminatory nucleic acid or composition is administerial an immunostiminatory nuclei temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomacosis, vitiligo, Crohn's Disease, ulcerative colitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated disetes mellitus, Grave's Disease, Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma,

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Gaps

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1; Indels

Mismatches

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16; Conservative

Matches

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The invention relates to an oligonuclectide having the formula: (a) 5'
XYN1YZN2 3' where 5' denotes the 5' end of the oligonuclectide and 3'
denotes the 3' end of the oligonuclectide, where X is a T or modified T
cuclectide, Y is a C or modified C nuclectide, Z is a G or modified G
nuclectide, and N1 and N2 are polynuclectides that do not include a CG
dinuclectide, where N1 does not include 5' Z nuclectide, and where 3'
colynuclectide consisting of the YZ dinuclectide and the N2
polynuclectide consisting of the YZ dinuclectide and the N2
c polynuclectide contains a number of nuclectides that is at most 45% of
the number of nuclectides in the oligonuclectide and (b) 5' XYN1YZN2 3'
where 5' denotes the 5' end of the oligonuclectide and 3' denotes the 3'
c end of the oligonuclectide, where X is a T or modified T nuclectide, Y is
a C or modified C nuclectide, where X is a G or modified muclectide, N1 is a
polynuclectide of 5-10 nuclectides, where N1 does not include a CG
dinuclectide, where N1 does not include 5' Z nuclectide, and where N2 is
a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a
pharmaceutical composition comprising the oligonuclectide in combination
if a therappeutic agent selected from chemcherapeutic agents.
                                                                                                                                                                                                                          ö
polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autolimune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with a therapeutic agent selected from chemotherapeutic agents, radiotherapeutic agents, monoclonal antibodies, and anticancer agents; (2) a method of specifically increasing interleukin (IL)-10 expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease, arthritis, systemic lupus erythematosus, multiple
sclerosis, Crohn's disease, or Type 1 diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New oligonucleotides, useful for treating an allergy or asthma, or an
                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.
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                                                                                                                                                                            Score 15.4; DB 1; Length 17; Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IL-10 expression assay, test oligonucleotide SEQ ID No:44.
                                                                                                                                                                                                                        1; Indels
                                                                                                                                Sequence 17 BP; 0 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                            Query Match
Best Local Similarity 94.1%;
Matches 16; Conservative
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                                                                                         invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1557
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call response to interferon (IFN)-alpha expression in a subject, comprising administering an oligonucleotide or a pharmaceutical composition to the subject in need of increased IL-10 expression relative to IFN-alpha expression; (3) a method of inducing an antigen specific regulatory T call response in a subject by administering an immunostimulatory nucleic acid or composition to a subject exposed to an antigen; (4) a method of inducing an antigen. Specific regulatory B cell response in a subject by administering an antigen. Specific regulatory nucleic acid or composition to a subject by administering an antigen. Specific acid or composition to the subject by astronaulatory nucleic acid or composition is administered in an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount subject by exposing a subject to a self antigen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount subject to a self antigen, and administered in an amount subject to a subject to an implant and administered in an amount sufficient to prevent or treat an autoimmune disease in the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigen administering an immunostimulatory nucleic acid or composition are useful for treating an action are useful for treating and composition are useful for treating allergies, asthma, autoimmune the invention are useful for treating allergies asthma, autoimmune the subject and a subject such as a current and as durable such as Guillain Barre, autoimmune their autoimmune their antigen are and a subject enemal, permic temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, vitiligo, Crohn's Disease, ulcerative colitis, primary biliary cirrhosis, autofimune hepatitis, immune-mediated diabetes mellitus, Grave's Disease. Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma, polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an Gaps used in experiments in the examples of the present pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss. ö Length 17; IL-10 expression assay, test oligonucleotide SEQ ID No:28. 1; Indels Sequence 17 BP; 0 A; 1 C; 1 G; 15 T; 0 U; 0 Other; Score 15.4; DB 1; Pred. No. 1.2e+03; 0; Mismatches 1; ° 2723 묤. Local Similarity 94.1%; es 16; Conservation Н 2707 CTAAAAAAAAAAAAAAA 17 CGAAAAAAAAAAAAAA AED81270 standard; DNA; 17 26-JAN-2006 (first entry) oligonucleotide WO2005111057-A2. 24-NOV-2005. invention Synthetic AED81270; Query Match RESULT 1558 Matches AED81270, BXBXGXEEXBXBXGXB ઠે 원

04-APR-2005; 2005WO-US011827.

02-APR-2004; 2004US-0558951P.

(COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH.

Krieg AM, Vollmer J;

WPI; 2005-786756/80.

New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus.

Example; SEQ ID NO 28; 111pp; English.

The invention relates to an oligonuclectide having the formula: (a) 5, XYNIYZNZ 3' where 5' denotes the 5' end of the oligonuclectide and 3' denotes the 3' end of the oligonuclectide, X is a T or modified G conclectide, Y is a C or modified G conclectide, and NI and NZ are polynuclectides that do not include a CC dinuclectide, where NI does not include 5' Z nucleotide, and where a 3' polynuclectide contains a number of mucleotide and the NZ polynuclectide contains a number of mucleotide and 3' denotes the 3' end of the oligonuclectide and 3' denotes the 3' cof the number of nucleotides in the oligonuclectide and 3' denotes the 3' cm of the oligonuclectide and 3' denotes the 3' cm of the oligonuclectide and 3' denotes the 3' cm of the oligonuclectide, where NI does not include a CC cm of the oligonuclectide, where NI does not include a CC cm oliveride of 5-10 nuclectides, where NI does not include a CC cm oliveride of 5-10 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a polynuclectide of 5 to 30 nuclectides. Also described are: (1) a captorherapeutic agents, monoclonal anticodies, and anticancer agents, radiotherapeutic agents, monoclonal anticodies, and anticancer agents, crelative to interferon (IFN) alpha expression in a subject, compression crelative to interferon (IFN) alpha expression in a subject or composition to the subject or an antigen. Specific regulatory B cell response in a subject by administering an antigen-specific acid or composition to a subject exposed to an antigen, specific regulatory B cell response in an antigen and administering an antigen acid or composition is administering an antigen acid or composition is admi Crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune neuropathies such as Guillain-Barre, autoimmune uveitis, autoimmune hemolytic anemia, pernicious anemia, autoimmune thrombocytopenia, temporal arteritis, anti-phospholipid syndrome, psoriaais, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, vitiligo, Crohn's blesses, ulcerative collitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated diabetes mellitus, Grave's Disease, Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune amount sufficient to prevent or treat an autoimmune disease in the subject, and (7) a method of reducing an antigen-specific response to an implant in a subject by exposing a subject to an implant and administering an immunostimulatory nucleic acid or composition to the in a subject by exposing a subject to a self antigen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an phosphorothicate linkage. The oligonuclectide, methods and compositions of the invention are useful for treating allergies, asthma, autoimmune diseases, arthritis, systemic lupus erythematosus, multiple sclerosis, subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigenspecific response to the implant in the subject. The oligonucleotide includes at least 1 modified internucleotide linkage such as a

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                     polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimume disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present
                                                                                                                                                                                                                             Gaps
  disease of the adrenal gland, rheumatoid arthritis, scleroderma,
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                                                                                                                                                                                   0.6%; Score 15.4; DB 1; Length 17; 44.1%; Pred. No. 1.2e+03;
                                                                                                                                                                                                                             1; Indels
                                                                                                                                          Sequence 17 BP; 1 A; 0 C; 0 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                    2709 AAAAAAAAAAAAAA 2725
                                                                                                                                                                                                      94.1%;
                                                                                                                                                                            Query Match
Best Local Similarity 94.15
Matches 16; Conservative
                                                                                                      invention.
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17 AAAAAAAAAAAAATA 1

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RESULT 1559

ВР AED81243 standard; DNA; 17 AED81243,

(first entry) 26-JAN-2006 AED81243;

IL-10 expression assay, test oligonucleotide SEQ ID No:1.

pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.

Synthetic.

WO2005111057-A2.

24-NOV-2005

04-APR-2005; 2005WO-US011827.

02-APR-2004; 2004US-0558951P.

(COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH.

Vollmer J; Krieg AM,

WPI; 2005-786756/80.

autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus. New oligonucleotides, useful for treating an allergy or asthma,

Example; SEQ ID NO 1; 111pp; English.

The invention relates to an oligonucleotide having the formula: (a) 5'

XYNIYZM2 3' where 5' denotes the 5' end of the oligonucleotide and 3'
denotes the 3' end of the oligonucleotide, where X is a T or modified T
nucleotide, Y is a C or modified C nucleotide, Z is a G or modified G
incleotide, where NI does not include 5' Z nucleotide, and where a 3'
polynucleotide consisting of the YZ dinucleotide and the NZ
polynucleotide contains a number of nucleotides that is at most 45% of
the number of nucleotides in the oligonucleotide, and (b) 5' XYNIYZN2 3'
where 5' denotes the 5' end of the oligonucleotide, and (b) 7' XYNIYZN2 3'
where 5' denotes the 5' end of the oligonucleotide and 3' denotes the 5'
end of the oligonucleotide, where X is a T or modified T nucleotide, Y is a
C or modified C nucleotide, and end of the oligonucleotide, NI is a
polynucleotide of 5-10 nucleotides, where NI does not include a CG
dinucleotide, where NI does not include 5' Z nucleotide, and where NZ is
a polynucleotide of 5 to 3 nucleotides. Also described are: (1) a
polynucleotide of 5 to 3 nucleotides. Also described in combination
with a therapeutic agent selected from chemotherapeutic agents;
radiotherapeutic agents, monoclonal antibodies, and anticancer agents;

allergen in the subject; (6) a method of treating an autoimmune disease allergen in the subject; (6) a method of treating an autoimmune disease in a subject by exposing a subject to a self antigen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or treat an autoimmune disease in the subject; and (7) a method of reducing an antigen-specific response to an implant in a subject by exposing a subject to an implant antigen, and administering an immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an amount sufficient to prevent or reduce an antigenspecific response to the implant in the subject. The oligonucleotide includes at least I modified internucleotide linkage such as a phosphorothicate linkage. The oligonucleotide, methods and compositions of the invention are useful for treating allergies, asthma, autoimmune diseases, arthritis, systemic lupus erythematosus, multiple solerosis, crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune neuropathies such as Guillain-Barre, autoimmune uveitis, autoimmune hemolytic anemia, pernicious anemia, autoimmune thrombocytopenia, relative to interferon (IRN) another superson in a subject, comprising administering an oligonucleotide or a pharmaceutical composition to the subject in need of increased IL-10 expression in a subject, comprising expression, (3) a method of inducing an antiqen-specific regulatory T cell response in a subject by administering an immunostimulatory T acid or composition to a subject exposed to an antigen; (4) a method of inducing an antigen; (4) a method of inducing an immunostimulatory nucleic acid or composition to a subject exposed to an antigen; (5) a method of treating an allergy or subject exposed to an antigen; (6) a method of treating an allergy or subject to an antigen; (7) a method of treating an allergy or immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition to the subject, where the immunostimulatory nucleic acid or composition is administered in an polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma, method of specifically increasing interleukin (IL)-10 expression temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, villigo, Crol Disesse, ulcerative colitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated diabetes mellitus, Grave's Disease, oligonucleotide used in experiments in the examples of the present 

Seguence 17 BP; 0 A; 1 C; 0 G; 16 T; 0 U; 0 Other;

Gaps , 0 0.6%; Score 15.4; DB 1; Length 17; 94.1%; Pred. No. 1.2e+03; Ive 0; Mismatches 1; Indels 94.1%; Best Local Similarity 94.1 Matches 16; Conservative Query Match

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IL-10 expression assay, test oligonucleotide SEQ ID No:29. AED81271 standard; DNA; 17 BP. 26-JAN-2006 (first entry) AED81271; RESULT 1560 

pharmaceutical; therapeutic; immune stimulation; immune response; allergy; asthma; autoimmune disease; antiasthmatic; antiallergic; immunosuppressive; phosphorothioate; ss.

Synthetic.

WO2005111057-A2.

24-NOV-2005.

04-APR-2005; 2005WO-US011827.

02-APR-2004; 2004US-0558951P.

(COLE-) COLEY PHARM GROUI (COLE-) COLEY PHARM GMBH

Krieg AM, Vollmer J;

WPI; 2005-786756/80.

New oligonucleotides, useful for treating an allergy or asthma, or an autoimmune disease, arthritis, systemic lupus erythematosus, multiple sclerosis, Crohn's disease, or Type 1 diabetes mellitus.

temporal arteritis, anti-phospholipid syndrome, psoriasis, pemphigus vulgaris, vasculitis such as Wegener's granulomatosis, vitiligo, Crohn's Disease, ulcerative colitis, primary biliary cirrhosis, autoimmune hepatitis, immune-mediated diabetes mellitus, Grave's Disease, Crohn's disease, Type I diabetes mellitus, myasthenia gravis, autoimmune neuropathies such as Guillain-Barre, autoimmune uveitis, autoimmune hemolytic anemia, pernicious anemia, autoimmune thrombocytopenia, Example, SEQ ID NO 29, 111pp; English.

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The invention describes a host cell transformed or transfected with a nucleic acid encoding a plant-derived carotenoid cleavage dioxygenase (CCD) enzyme. Also described are: (1) a plasmid or vector system comprising a nucleotide including one of 1617, 1696, 1653, or 1557-amino acid sequence (SEQ ID No. 1, 3, 5, or 9) given in the specification or a sequence that is 754 homologous; (2) producing (M1) a carotenoid cleavage compound comprising treating a carotenoid with a plant-derived CCD; (3) an enzyme comprising the amino acid sequence corresponding to Rubus an enzyme comprising the amino acid sequence corresponding to Rubus isolated nucleic acid molecule coding for the enzyme, comprising a nucleotide sequence that is the same as, or complementary to a 2429 base nucleotide sequence that is the same as, or complementary to a 2429 base pair sequence (SEQ ID NO.3) and (5) a CrtL-e enzyme comprising 527-amino acid sequence (SEQ ID No. 10) or its effective fragment. The host cell is used for producing a carotenoid cleavage compound, e.g. alpha or beta ionone, pseudo ionone, safranal, theaspirone, damascone or damascenome, and for producing GGDP, beta -carotene, lycopene, or delta carotene for use in flavoring or perfume applications including e.g. soft drinks, fruit juice or beverage
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Hashimoto's thyroiditis, autoimmune oophoritis and orchitis, autoimmune disease of the adrenal gland, rheumatoid arthritis, scleroderma, polymyositis, dermatomyositis, spondyloarthropathies such as ankylosing spondylitis, and Sjogren's syndrome. The autoimmune disease is caused by an infection e.g. Lyme disease. This sequence represents an oligonucleotide used in experiments in the examples of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transformation; genetically engineered microorganism; PCR; primer; ss.
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                                                                                                                                                                                                                                                               Gaps
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nucleic acid encoding plant-derived carotenoid cleavage deoxygenase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Host cell for producing carotenoid cleavage compound for flavoring
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                                                                                                                                                                                                              Score 15.4; DB 1; Length 17;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Carotenoid cleavage dioxygenase associated primer PolyT.
                                                                                                                                                                        Sequence 17 BP; 0 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                       2709 AAAAAAAAAAAAAA 2725
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEF63888 standard; DNA; 17 BP
                                                                                                                                                                                                                Query Match

Best Local Similarity 94.1%;
Matches 16; Conservative C
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                                                                                                                                   invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ovist I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Poly T primer, to synthesize red raspberry CHS cDNA and rhubarb BAS cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Host cell comprising a benzalacetone synthase (BAS) polypeptide sequence or a 4-coumarate:CoA ligase (4CL) sequence, useful in a bacterial method of producing benzalacetone and/or raspberry ketone is new.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Flavor enhancer; food; cosmetics; weight loss; naringenin-chalcone synthase;
                                                                                                                                                                                                                                                         techniques. This sequence represents a primer associated with the isolation and cloning of DNA encoding a carotenoid cleavage dioxygenase
comprising whey protein, health teas, cocoa drinks, milk drinks and lactic acid bacterial drinks, yogurt, drinking yogurt and wine, bakery product including bread, Danish pastry, biscuits or cookies, confectionery product, pharmaceutical composition for therapeutic or diagnostic purposes. The invention provides reliable and efficient production of aroma compounds or precursors, particularly carotemoid cleavage compounds that does not rely solely on chemical synthesis techniques. This sequence represents a primer associated with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hall RD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 15.4; DB 1;
18.2%; Pred. No. 1.2e+03;
ve 2; Mismatches 0;
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AEF88073/
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The present invention relates to a host cell comprising a chalcone synthase (GHS; also referred as naringenin-chalcone synthase, flavanone synthase; 6'-deoxychalcone synthase; flavanone synthase; 6'-deoxychalcone synthase; chalcone synthetase and DOCS; EC 2.3.1.74) polypeptide sequence and 4-coumarate.CoA ligase (4CL) also referred as 4-coumarcyl-CoA synthetase; p-coumaryl coenzyme A synthetase; p-coumaryl CoA synthetase; p-coumaryl-CoA synthetase; p-coumaryl-CoA synthetase; feruloyl Coenzyme A synthetase; p-coumaryl-CoA synthet
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production of benzalacetones and/or raspberry ketones from p-coumaric acid. Raspberry ketone is a flavor component used in the food industry. Benzalacetones and raspberry ketones are useful in aroma formulations in food market, cosmetics, household products such as air fresheners and in weight loss products. The present sequence is a PCR primar used in the synthesis of red raspberry chalcone synthase (CHS) cDNA and benzalacetone synthase (BAS) cDNA, which is used in the production of raspberry ketone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Host cell used in producing benzalacetone or raspberry ketone comprises chalcone synthase polypeptide sequence and 4-coumarate:COA ligase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Poly T primer used to synthesize red raspberry chalcone synthase (CHS).
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                                                                                                                                                                                                                                                                                                                                                    Gaps
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naringenin-chalcone synthase; chalcone synthase; flavanone synthase;
6'-deoxychalcone synthase; chalcone synthetase; DOCS; primer; ss.
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                                                                                                                                                                                                                                                                              y Match 0.6%; Score 15.4; DB 1; Length 17; Local Similarity 88.2%; Pred. No. 1.2e+03; hee 15; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                 Seguence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beekwilder MJ, Sibbesen O, Mikkelsen JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 43; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rubus idaeus; cultivar Tulameen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 ВВАААААААААААААА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEF87959 standard; DNA; 17
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Matches
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cosmetics, household products such as air fresheners and in weight loss products. Chalcone synthase has benzalacetone synthase (BAS) activity. The present sequence is a primer used in the synthesis of red raspberry chalcone synthase (CHS) cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bind to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New sequence-specific non-photo-activated crosslinking agents - bind to
the major groove of duplex DNA and are esp. useful for treating latent
infections e.g. HIV.
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                                                                                                                                                                                                                                                                                                                                                        deoxyribonucleic acid; major groove; ethanoamino group; tumour necrosis factor; receptor; messenger RNA; aziridinylcytosine; cross-linking group; ss.
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                                                                                                                                                                                                                                                                                                                                 Cross-linking oligomer 943 to target human TNF Receptor mRNA.
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                                                                                               Score 15.4; DB 1; Length 17;
Pred. No. 1.2e+03;
2; Mismatches 0; Indels
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/mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
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                                                                       Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
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/note= "N4N4-ethanocytosine"
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                                                                                                                             15; Conservative
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                                                                                                   Query Match
Best Local Similarity
Matches 15; Conserv
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modified_base
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RESULT 1565

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AAQ20108/

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Sequence beginning at mucleotide 2354. The sequence is a viral duplex sequence beginning at mucleotide 2354. The sequence is a viral duplex sequence contg. a purine-rich region concentrated on one chain of the duplex. The sequence may be prepd. by standard DNA synthesis. The HUMNFR of forming a triplex at physiological pH by coupling into the major groove of the DNA duplex. Three such oligomers which are capable of forming a triplex with this sequence. The oligomers are used in the treatment of inflammation. Similar oligomers may be used to target viral DNA duplexes specific for HIV, herpes and other viruses. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also
                                                                           Target; human tumour necrosis factor receptor mRNA; AIDS; triplex; HIV;
hepatitis; malignancy; inflammation; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence depicts a HUMNFR (tumour necrosis factor receptor) mRNA
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                                      Purine rich HUMNFR target duplex sequence.
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
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     (first entry)
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(first entry)
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18-JAN-1991;
08-APR-1991;
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17-APR-1991;
17-APR-1991;
     07-DEC-1992
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07-DEC-1992
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                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                           deoxyribonucleic acid; major groove; ethanoamino group;
tumour necrosis factor; receptor; messenger RNA; aziridinylcytosine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "N4N4-ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mod_base= OTHER
2709 AAAAAAAAAAAAAA 2725
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/mod_base= m5c
                                                                                                                            AAQ20108 standard; DNA; 18 BP
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                                17 AAAAAAAAAAATAAA 1
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                                                                                                                                                                                                                                                                                                                  cross-linking group; ss.
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14-JAN-1991;
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AAQ25501;

RESULT 1566

AAQ25501

Query Match

Best Loca Matches

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Human tumour necrosis factor receptor mRNA; AIDS; modified; HIV; RSV;
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                                                                                                  '*tag= a
mod base= OTHER
note= "N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                          Matteucci MD, Milligan J;
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/note= "OTHER= N4 N4 ethanocytosine'
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                HPV; malignancy; hepatitis; inflammation; ss.
                                                                    Location/Qualifiers
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91US-00686546.
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modified_base
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17-APR-1991;
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07-DEC-1992
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                                           Synthetic
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The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor receptor mRNA beginning at nucleotide 2134 contg, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HER, HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PD field.)
                                 Human tumour necrosis factor receptor mRNA; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; 88.
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        Oligomer TNFR942 for forming triplex with HUMNFR target duplex.
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                                                                                                                                                                                            /mod_base= OTHER
/note= "OTHER= N4 N4 ethanocytosine"
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                                                                                                         location/Qualifiers
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910S-00683420.
910S-00686544.
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910S-006865473.
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Best Local Similarity
                                                                                                              Key
modified_base
                                                                                                                                                                 modified_base
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08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
27-SEP-1991;
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                                                                                 Synthetic.
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AAV54168/C
ID AAV54168
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AC AAV54168
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This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis of a (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases
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                                       Novel apoptosis-related DNAs and proteins - for diagnosis, preventing or treating diseases associated with apoptosis.
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                                                                                                Example 1; Page 47; 70pp; Japanese.
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ID AAV54166 standard; cDNA; 18
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Best Local Similarity 94.1
Matches 16; Conservative
      WPI; 1998-495844/42.
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                                       Nucleotide sequence PCR primer 5.
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(first entry)
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This is the nucleotide sequence of a PCR primer used in the method of the invention, involving the use of novel apoptosis-related DNAs and proteins. The inventions can be used as diagnostic reagents for apoptosis e.g. (monoclonal) antibodies for the protein, as a reagent in immunohistological staining, as apoptosis inhibitors. It can also be used for treatment of apoptosis-related diseases
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Local Similarity 94.1%; Pred. No. 1.2e+03;
les 16; Conservative 0; Mismatches 1; Indels
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AAV54169 standard; cDNA; 18
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                                            Homo sapiens.
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jical staining, as apoptosis inhibitors. It can also be used of apoptosis-related diseases
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                                                0.6%; Score 15.4; DB 1; Length 18; 94.1%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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                               BP; 0 A; 0 C; 3 G; 15 T; 0 U; 0 Other;
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                                                                                                                                                                                                                     Human adipose tissue gene amplifying primer #9.
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                                                                                      2707 CTAAAAAAAAAAAA 2723
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                                                                     16; Conservative
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   immunohistological
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             for treatment
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Adipose tissue; obesity; diabetes; hyperlipemia; hypertension; human; arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss.
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Best Local Similarity 94.19
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tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperuticemia and sleep apnea syndrome. The genes (AAZ90631-633) and the proteins (AAX67598-Y67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose tissue genes
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Best Local Similarity 94.1%; Pred. No. 1.2e+03;
Matches 16; Conservative 0; Mismatches 1; Indels
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                                                                                Sequence 18 BP; 0 A; 1 C; 2 G; 15 T; 0 U; 0 Other;
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                                                    Adipose tissue, obesity; diabetes; hyperlipemia; hypertension; human; arteriosclerosis; hyperuricemia; sleep apnea syndrome; PCR primer; ss.
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Human adipose tissue gene amplifying primer #6.
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Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                        23-JUL-1998;
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                                                                                                                                                    Homo sapiens
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Example 2; Page 18; 50pp; Japanese.
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The invention relates to identification of genes and proteins of adipose tissue relating to obesity, particularly complications of visceral obesity including diabetes, hyperlipemia, hypertension, arteriosclerosis, hyperuricemia and sleep apnea syndrome. The genes (AAZ90631-633) and the proteins (AAZ6759B-Y67600) are used in the genetic diagnosis, prevention and treatment of adipose tissue related diseases. Sequences AAZ90640-51 represent PCR primers amplifying the human adipose tissue genes

Sequence 18 BP; 1 A; 0 C; 2 G; 15 T; 0 U; 0 Other;

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Score 15.4; DB 1; Length 18; Pred. No. 1.2e+03;
                    Indels
                    Mismatches
                                       2724
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0.6%;
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                                      2708 TAAAAAAAAAAAA
                  Conservative
         Local Similarity
                  16;
 Query Match
                 Matches
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Gaps

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.554/c AAZ70554 standard; DNA; 18 BP. (first entry) 10-SEP-2001 AAZ70554; RESULT 1581 

Human biallelic marker upstream amplification primer SEQ ID NO:4910. Human genome, biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;

diagnosis; ss

Homo sapiens

WO9954500-A2

28-OCT-1999

99WO-IB000822 21-APR-1999; 98US-0082614P, 98US-0109732P, 23-NOV-1998; 21-APR-1998;

(GEST ) GENSET

Chumakov I; Blumenfeld M, Cohen D,

WPI; 2000-013267/01

Novel biallelic markers used to construct a high density disequilibrium map of the human genome

Claim 8; Page 1278; 2745pp; English.

AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ6579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and

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3367, are not actually given a sequence in the Sequence Listing from the
                                    Gaps
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0
                          Length 18;
                                    1; Indels
               Sequence 18 BP; 0 A; 8 C; 1 G; 9 T; 0 U; 0 Other;
                        Score 15.4; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 1;
                                                                                                                SNP specific lower PCR primer SEQ ID 710.
                                    o;
                                              717
                                                                                  BP.
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                         0.6%;
                                             701 GCAGAGGAAGAACAAGA
                                                                                AAH37914 standard; DNA; 18
                                                        GGAGGGAAGAACAAGA
                                                                                                     (first entry)
                                    16; Conservative
                              Similarity
      present invention
                                                                                                      14-AUG-2001
                                                                                           AAH37914;
                                                        18
                         Query Match
Best Local
                                                                       RESULT 1582
                                  Matches
                                                                             AAH37914/
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Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple solerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss. Homo sapiens

WO200129262-A2

26-APR-2001.

13-OCT-2000; 2000WO-US028436.

99US-0160096P. 15-OCT-1999; (ORCH-) ORCHID BIOSCIENCES INC

Pohl M; Picoult-Newburg L,

WPI; 2001-290930/30.

New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic acid sample.

Claim 1; Page 53; 83pp; English.

primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the coligonucleotides of the invention. The PCR primers are used to amplify a cliquance of the invention. The PCR primer are used to amplify a coligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for genotyping a nucleic acid sample by performing a single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or cliqual single-nucleotide primer extension reaction. The oligonucleotides are useful for determining the presence, absence or cliqual shaving a pathological phenotypic trait suspected of being assess by association analysis the genotype of an individual or group of individuals, having a pathological phenotypic traits suspected of being caused by one or more SNPs. Phenotypic traits include diseases e.g. caused by one or more SNPs. Phenotypic traits include disease e.g. dystrophy, familial hypercholesterolemia, polycystic kidney disease, osteogenesis imperfecta and acute intermittent porphyria. Phenotypic traits also include symptoms of or susceptibility to multifactorial disease of which a component is or may be genetic such as autoimmune diseases, including, rheumatoid arkhitis, multiple sclerosis, inflammation, cancer, nervous system diseases and infection by pathogenic Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide

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RESULT 1584
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microorganism. The method is also useful in forensic investigations and paternity analysis. The present sequence represents a PCR primer specific for a human SNP containing DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        which are capable of metabolizing the drug. A method for treatment of tachyphylaxis therefore involves inhibiting the induced metabolic enzyme,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a metabolic
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14.1%; Pred. No. 1.2e+03;
ve 0; Mismatches 1; Indels
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                                                                                                                                                    Sequence 18 BP; 3 A; 6 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                      18 GCCCTGAAACTTGAGG 2
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Best Local Similarity 94.1%;
Matches 16; Conservative
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DB 1; Length 18;

0.6%; Score 15.4;

Query Match

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The invention relates to an antisense oligonucleotide or its active derivative selected from AEC46374-AEC46395, targeting human interleukince derivative selected from AEC46374-AEC46395, targeting human interleukince derivative selected from AEC46374-AEC46395, targeting human interleukince oligonucleotide (or its active derivative, by adding consecutive coligonucleotide (or its active derivative, by adding consecutive concleotide (and interpretains), a pharmaceutical composition comprising concleotide chain), a pharmaceutical composition compraising composition for treating cancer. The oligonucleotide is an antisense composition for treating cancer. The oligonucleotide is an antisense coligonucleotide inhibiting the synthesis of proteins involved in the coligonucleotide inhibiting the production of transforming growth factor (TGF)-beta 1, TGF-beta 2, TGF-beta 3, cell-cell adhesion molecules coligonucleotide inhibiting the production of transforming growth factor (TGF)-beta 1, TGF-beta 2, TGF-beta 3, cell-cell adhesion molecules colimbitions (THPS) and/or interleukins 10. The oligonucleotides are useful for the preparation of a pharmaceutical composition for inhibiting cancer, breast cancer, bronchogenic Cancer, bronchogenic Cancer, bronchogenic Cancer, pastric cancer, color carcinoma, modullary carcinoma, non-small cell cancer, dastric cancer, head and neck cancer, hepatocellular cancer, cancer, head and neck cancer, hepatocellular carcinoma, color carcinoma, papillary adenocarcinoma, prostate cancer, small cell carcinoma, skin cancer, soft tissue cancer, squamous cell carcinoma, intestine carcinoma, skin cancer, soft tissue cancer
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                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides inhibiting the synthesis of proteins involved in the formation of metastases such as transforming growth factor-beta 1 (TGF-beta 1), TGF-beta 2 and TGF-beta 3, useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                       Transforming growth factor beta; TGF-beta-3; antisense therapy; antisense oligonucleotide; ss; cancer; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide targeting human TGF-beta-3 #1244.
                          Indels
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                        Mismatches
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    Pred. No.
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Bischof A, Hafner M, Egger T;
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                                                                                                                 1 AGCAGAGGAAGAGCAAG 17
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01-APR-2004; 2004US-0558135P.
  l Similarity 94.1%;
16; Conservative
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                                                                        700 AGCAGAGGAAGAACAAG
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Best Local Similarity
Matches 16; Conserv
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derivative selected from AEC46374-AEC46395, targeting human interleuking derivative selected from AEC46374-AEC46395, targeting human interleuking colliquoucleotide (or its active derivative, by adding consecutive nucleosides and linker stepwise or by cutting the oligonucleotide out of longer oligonucleotide chain), a pharmaceutical composition comprising the artisense oligonucleotide and a TGF-beta 2 antagonist for preparing a composition for treating cancer. The oligonucleotide is an antisense oligonucleotide inhibiting the synthesis of proteins involved in the composition of metastases. The oligonucleotide is an antisense oligonucleotide inhibiting the production of transforming growth factor (TGF)-beta 1, TGF-beta 3, Cell-cell adhesion molecules (CAMS), integrins, selectins, metalloproteases (MMPS), their tissue inhibitors (TIMPS) and/or interleukins 10. The oligonucleotides are useful for the preparation of a pharmaceutical composition for inhibiting the formation of metastases in cancer treatment. The oligonucleotides are
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testicular carcinoma, uterine cancer, acoustic neuromas, neurofibromas, trachomas, and pyogenic granulomas; pre-malignant tumors, blastoma, Ewing's tumor, craniopharyngioma, ependymona, medulloblastoma, hemangioblastoma, medullablastoma, etanoma, mesothelioma, neuroblastoma, hemangioblastoma, etinoblastoma, estroma, pinaloma, neuroblastoma, wilm's tumor and/or myeloma, multiple. The present sequence is an antisense oligonucleotide targeting human TGF-beta-3.
                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides inhibiting the synthesis of proteins involved in the formation of metastases such as transforming growth factor-beta 1 (TGF-beta 1), TGF-beta 2 and TGF-beta 3, useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide targeting human TGF-beta-3 #1104.
                                                                                                                                                                                                    DB 1; Length 18;
                                                                                                                                                                                                                                         Indels
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                                                                                                                                                           Sequence 18 BP; 5 A; 7 C; 6 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                  Score 15.4; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 1
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01-APR-2004; 2004US-0558135P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Best Local Similarity 94.1
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating cancer.
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               brain tumor, breast cancer, bronchogenic carcinoma, carcinoma of the kidney, cervical cancer, choriocarcinoma, cystadenocarcinoma, carvical carcinoma, colorectinoma, encrinoma, encrinoma, encrinoma, encrinoma, encrinoma, encrinoma, encrinoma, esophageal cancer, gall bladder cancer, gastric cancer, head and neck cancer, hepatocellular cancer, liver carcinoma, lung carcinoma, medullary carcinoma, non-small cell bronchogenic/lung carcinoma, ovarian cancer, pancreas carcinoma, papillary acroinoma, popillary adenocarcinoma, prostate cancer, small intestine carcinoma, rectal cancer, renal cell carcinoma, skin cancer, soft tissue cancer, squamous cell carcinoma, testicular carcinoma, uterine cancer, acoustic neuromas, neurofibromas, testicular carcinoma, uterine cancer, acoustic neuromas, neurofibromas, trachomas, and pyogenic granulomas; pre-malignant tumors, blastoma, Ewing's tumor, craniopharyngioma, ependymoma, medulloblastoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel hypoallergenic variants of group 1 allergens of Poaceae that have reduced immunoglobulin E (TgB) reactivitiy compared with the wild-type allergens but essentially retained reactivity with T lymphocytes. The invention also describes; a) DNA molecules that
                                                                                                                                                                                                                                                                                          hemangioblastoma, medullablastoma, melanoma, mesothelioma, neuroblastoma, eleuzofibroma, pintaaloma, retinoblastoma, aarcoma, seminoma, trachomas, Wilm's tumor and/or myeloma, multiple. The present sequence is an antisense oligonucleotide targeting human TGF-beta-3.
  useful for treating cancer, e.g. bile duct carcinoma, bladder carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New variants of group 1 grass pollen allergens, useful for treatment or prevention of allergy, have reduced immunoglobulin E reactivity while retaining T cell reactivity, also new DNA encoding them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; Phl p 1; allergen; immunoglobulin E; t-lymphocyte; vaccine; veterinary; Antiallergic; Hyposensitization; mutagenesis; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encode the variant allergens; b) a recombinant expression vector containing DNA encoding the variant allergens functionally linked to expression control sequence; c) a host organism transformed with DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                             0.6%; Score 15.4; DB 1; Length 18; 94.1%; Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 5 A; 6 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAGCAGGCCAGGACCA 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94.18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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Human chromosome 19q r region SNP containing region primer #22. encoding the variant allergens or the expression vector and d) 0.6%; Score 15.4; DB 1; Length 18; 4.1%; Pred. No. 1.2e+03; 1; Indels Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 U; 0 Other; 0; Mismatches (ARBE-) ARBEJDSMILJOINSTITUTTET Borglum A; 381 GGCGGGACCTCCGGGAT 397 BP. 23-FEB-2005; 2005DK-00000274. 22-JUN-2005; 2005DK-00000918. 17 decedencerredent 1 18-AUG-2004; 2004DK-00001249 17-AUG-2005; 2005WO-DK000529 94.18; AEF93735 standard; DNA; 18 (first entry) Matches 16, Conservative Vogel UB, WPI; 2006-173654/18 UYAA-) UNIV AARHUS Best Local Similarity WO2006018023-A2. Homo sapiens 20-APR-2006 23-FEB-2006 Nexo BA, AEF93735 Query Match AEF93735 원 ઠે

The invention relates to a method of estimating the disease risk of an individual which comprises assessing in the genetic material a sequence polymorphism, obtaining a sequence polymorphism response, and estimating the disease risk of the individual based on the sequence polymorphism response. The methods are useful for estimating the disease risk of an individual comprises, estimating the disease prognosis of an individual and estimating a treatment response of an individual suffering from cancer to a disease treatment. The methods can be used for identifying subjects with increased risk of having or developing cancer. The present sequence represents a human chromosome 19q r region PCR primer. The invention relates to a method of estimating the disease risk of an individual which comprises assessing in the genetic material a sequence polymorphism, obtaining a sequence polymorphism response, and estimating the disease risk of the individual based on the sequence polymorphism response. The methods are useful for estimating the disease risk of an individual comprises, estimating the disease prognosis of an individual and estimating a treatment response of an individual suffering from cancer to a disease treatment. The methods can be used for identifying subjects with increased risk of having or developing cancer. The present sequence represents a human chromosome 19q r region primer/probe. Bstimating disease risk of an individual, by assessing a sequence polymorphism, obtaining a sequence polymorphism response and estimating disease risk of individual based on the sequence polymorphism response. detection; prognosis; chromosome 19; r region; ss; probe; primer; Human chromosome 19q r region SNP containing region oligo #58. 0.6%; Score 15.4; DB 1; Length 18; 94.1%; Pred. No. 1.2e+03; ve 0; Mismatches 1; Indels Sequence 18 BP; 4 A; 9 C; 1 G; 4 T; 0 U; 0 Other; Claim 60; Page 98; 139pp; English. Example; Page 68; 139pp; English. (ARBE-) ARBEJDSMILJOINSTITUTTET 1825 GGAGAAGGAGGTTGCAG 1841 Borglum A; 18 GGAGATGGAGGTTGCAG 2 18-AUG-2004; 2004DK-00001249. 23-FEB-2005; 2005DK-00000274. 22-JUN-2005; 2005DK-0000091B. 17-AUG-2005; 2005WO-DK000529. 94.18; AEF93897 standard; DNA; 18 (first entry) 16; Conservative Vogel UB, WPI; 2006-173654/18 UYAA-) UNIV AARHUS Local Similarity WO2006018023-A2. Homo sapiens. 20-APR-2006 23-FEB-2006. Query Match AEF93897 Nexo BA, 1588 Matches RESULT 15 SNP ò В ö Betimating disease risk of an individual, by assessing a sequence polymorphism, obtaining a sequence polymorphism response and estimating disease risk of individual based on the sequence polymorphism response. Gaps SNP detection; prognosis; chromosome 19; r region; ss; PCR; primer. . 0

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Unidentified
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Createsion (SNPE) primers, and the sequences of regions flanking

Sites of single nucleotide polymorphisms SNPs. The present invention

includes kits for determining the presence or absence of a SNP, using the

includes kits for determining the presence or absence of a SNP, using the

Creates are useful for genotyping a nucleic acid sample by

Derforming a single-nucleotide primer extension reaction. The

Cligonucleotides are useful for determining the presence, absence or

identity of a SNP and for genotype of an individual or group of

clidentity of a SNP and for genotype of an individual or group of

individuals, having a pathological phenotypic trait suspected of being

caused by one or more SNPs. Phenotypic trait suspected of being

caused by one or more SNPs. Phenotypic trait suspected of being

caused by one or more SNPs. Phenotypic traits include diseases e.g.

agammaglobulinaemia, diabetes insipidus, Lesch-Nyhan syndrome, muscular

dystrophy, familial hypercholesterolaemia, polycystic kidney disease,

cateogenesis imperfecta and acute intermittent porphyria. Phenotypic

ctraits also include symptoms of or susceptibility to multifactorial

cdisease of which a component is or may be genetic such as autoimmune

diseases, including, rheumatoid arthritis, multiple sclerosis,

microorganism. The method is also useful in forensic investigations and

paternity analysis. The present sequence

con a human SNP containing DNA sequence
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                                                                                                                                                                                                                                                                                                               Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-NWhan syndrome; nuscular dyterophy; familial hypercholesteroleamia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nucleic
                                                            Gaps
                                                            ;
                             Score 15.4; DB 1; Length 18; Pred. No. 1.2e+03;
                                                            Indels
Sequence 18 BP; 4 A; 9 C; 1 G; 4 T; 0 U; 0 Other;
                                                        0; Mismatches
                                                                                                                                                                                                                                                                                      SNP specific lower PCR primer SEQ ID 3718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 68; 83pp; English.
                                                                                       1825 GGAGAAGGAGGTTGCAG 1841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ORCH-) ORCHID BIOSCIENCES INC.
                                                                                                                                                                                            AAH40922 standard; DNA; 19 BP.
                             0.6%;
                                                                                                             18 GGAGATGGAGGTTGCAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-OCT-2000; 2000WO-US028436
                                                                                                                                                                                                                                                      14-AUG-2001 (first entry)
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-290930/30.
                     Query Match
Best Local Similarity
Matches 16; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acid sample.
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                                                                                                                                                                                                                          AAH40922;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention comprises primers for analysing the expression of cytochrome P450 1 Al gene and cytochrome P450 Bl0 gene, and estimating the toxicity of a test agent. The PCR primers of the invention are useful for analysing tan expression of cytochrome P450 1 Al gene and P450 2Bl0 gene and estimating the toxicity of a test agent. The present DNA sequence represents a PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligonucleotide useful for analyzing expression of cytochrome P450 1 Al gene and cytochrome P450 BlO gene, and estimating toxicity of test
                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome P450 B10; toxicity estimation; PCR;
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14.1%; Pred. No. 1.3e+03;
ve 0; Mismatches 1; Indels
                                                   0.6%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.3e+03; ve 0; Mismatches 1; Indels
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Sequence 19 BP; 4 A; 2 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytochrome P450 gene-specific PCR primer #14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer; expression analysis; ss
                                                                                                                                                                2460 CCCTCACCAAGCACTTC 2476
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                                                                                Best Local Similarity 94.1
Matches 16; Conservative
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les 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome P450
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polycystic kidney disease; RNA interference; short interfering RNA; siRNA; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene thrapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cytostatic; vasotropic;

nephrotropic; ss

WO2003070917-A2. Homo sapiens.

28-AUG-2003

Human; Myc; Myb; cancer; proliferative disease; restenosis;

Human Myc siNA lower strand, SEQ ID 192

12-FEB-2004 (first entry)

ADF48055;

ADF48055 standard; RNA; 19 BP

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human Myc or Myb genes by RNA interference. The sinAs may or comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an misense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double specifically, the siNAs include short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the Myc or Myb genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and used for treating cancers and other proliferative diseases, such as retenosis and polycystic kidney disease. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human Myc-targeted challed and and bene mapping the upper strand of a human Myc-targeted challed polymorphisms).
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Human; Myc; Myb; cancer; proliferative disease; restenosis; polycystic kidney disease; RNA interference; short interfering mucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene function analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 9 A; 1 C; 8 G; 0 T; 1 U; 0 Other;
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2002US-0418655P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-689784/65.
                                                                                                                                                                                                                                                                                                                                       nephrotropic; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003070917-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                            Ното варієпв.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2002;
11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
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2002US-0408378P. 2002US-0409293P. 2002US-0418655P.

05-SEP-2002; 09-SEP-2002; L5-JAN-2003;

2003US-0440129P

(RIBO-) RIBOZYME PHARM INC Mcswiggen J, Beigelman L;

WPI; 2003-689784/65.

2002US-0406784P

20-FEB-2003; 2003WO-US005326

2002US-0358580P 2002US-0363124P 2002US-0386782P

> 11-MAR-2002; 06-JUN-2002; 29-AUG-2002;

20-FEB-2002;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human Myc or Myb genes by RNA convergulate expression of the human Myc or myb genes by RNA convergulate expression of the human Myc or myb genes by RNA controlled for an antisense and antisense and antisense or single stranded. They further comprise sense and antisense coligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Cc an contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express sinA. The siNAs are used to modulate expression of the Myc or Myb genes in cells, tissue cy transplants for the treatment of a variety of conditions. They may be consecuted in conditions. They may be consecuted in the conditions and consecution and consist and polycystic kidney disease. The siNAs are also useful for drug screening, genetic engineering, pharmacogenomics, studying gene condition, and gene mapping (e.g., of single nucleotide polymorphisms).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 1 A; 8 C; 1 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double-stranded siNA.
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New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of Myc or Myb genes.

Example 7; Page 128; 161pp; English

Length 19; Score 15.4; DB 1; Pred. No. 1.3e+03; 0.6%; Query Match Best Local Similarity

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Gaps

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0.6%; Score 15.4; DB 1; Length 19; 14.1%; Pred. No. 1.3e+03; ve 0; Mismatches 1; Indels

704 GAGGAAGAACAAGAAGA 720

ð g

Local Similarity 94.1%; les 16; Conservative

Query Match

Best Loca Matches

1 GAGGAGGAACAAGAAGA 17

Matches

RESULT 1593

g ò

ADL253

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SARS; severe acute respiratory syndrome; siRNA; short interfering RNA; ss; RNA interference; gene silencing; SARS virus infection; acute respiratory failure; viral pneumonia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acid obtained from 4 synthesis-period of neuroblastoma cells useful for prognosing and determining progress stage of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to human nucleic acid sequences (I; AD015739-AD015912) obtained from 4 synthesis-period (stage 4S) of neuroblastoma cell. (I) is useful for prognosing and determining th progress stage of 4 synthesis-period of neuroblastoma. The present sequence is a primer, used to illustrate the invention.
                                                                                               4 synthesis-period of neuroblastoma related primer, SEQ ID 781.
                                                                                                                                    Human; 4 synthesis-period; neuroblastoma; stage 4S; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.6%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.30+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 8 A; 6 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SARS coronavirus siRNA lower sequence 177.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 8; SEQ ID NO 781; 455pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGCTGTGGTGTTCTAT 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                      30-OCT-2003; 2003WO-JP013932.
                                                                                                                                                                                                                                                                                                                       30-OCT-2002; 2002JP-00316586.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-APR-2003; 2003US-0462874P. 30-APR-2003; 2003US-00427160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-APR-2004; 2004WO-US011320.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADT64925 standard; RNA; 19
                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 94.1
                                                                                                                                                                                                                                                                                                                                                                               (CHIB-) CHIBA PREFECTURE
                                                                                                                                                                                                                                                                                                                                                              PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                      Ohira M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-390323/36.
                                                                                                                                                                                                                                                                                                                                                              (HISM ) HISAMITSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SARS coronavirus
                                                                                                                                                                                                          WO2004039975-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004092383-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neuroblastomas,
                                                                                                                                                                                                                                                                                                                                                                                                                      Nakagawara A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JAN-2005
                                                            29-JUL-2004
                                                                                                                                                                                                                                                 13-MAY-2004.
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                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       903
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises DNA sequences which are associated with intestinal epithelium and peyer's patch M cells. The DNA sequences of the invention are useful for assessing, modifying, modulating or regulating intestinal epithelium or M cell development. The DNA sequences of the invention are also useful in the treatment of: inflammatory bowel disease, glutenenteropathy, infectious diseases, autoimmune diseases diseases, multiple sclerosis, allergy, asthma and diabetic mellitus), diseases or disorders of the immune system, hypersensitivity, anaphylaxis, and blood group incompatibility. The present DNA sequence represents, a PCR primer that was used to amplify an intestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                               intestinal epithelium cell development; peyer's patch M cell development; inflammatory bowel disease; glutenenteropathy; infectious disease; autoimmune disease; haemolytic anaemia; rheumatoid arthritis; dermatitis; Grave's disease; multiple sclerosis; allergy; asthma; diabetic mellitus; immune system disorder; hypersensitivity; anaphylaxis; blood group incompatibility; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated or purified polypeptide encoded by genes associated with intestinal epithelium or M cell development, differentiation or function, useful for treating autoimmune diseases and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epithelium/peyer's patch M cell-associated DNA sequence of the invention.
                                                                                                                                                                                                                                                                              Intestinal epithelium/peyer's patch M cell-associated PCR primer #480.
     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Indels
     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mah SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 4 A; 10 C; 1 G; 4 T; 0 U; 0 Other;
   1;
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   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure, SEQ ID NO 845, 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (DIGI-) DIGITAL GENE TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O'mahony DJ,
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                                      720
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-APR-2001; 2001US-0281416P
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                                                                       19 GAGGAGGAACAAGAAGA
                                                                                                                                                                   ADL25335 standard; DNA; 19
                                    704 GAGGAAGAACAAGAAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADO16519 standard; DNA; 19
                                                                                                                                                                                                                                           (first entry)
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Best Local Similarity 94.1
Matches 16; Conservative
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Byrne D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-075470/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Macaca fascicularis
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                                                                                                                                                                                                                                           20-MAY-2004
16;
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                                                                                                                                                                                                      ADL25335;
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RESULT 1594

ઠે 요 AD016519/ ID AD01

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Gaps

2003US-0462874P. 2003US-00427160. 2003US-00444853.

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15-APR-2003;
                               30-APR-2003;
23-MAY-2003;
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 BXBX8X£4XBXBXBXB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          셤
                                                                                                                                                                                                                                                   The invention relates to a chemically synthesised double stranded short interfering ribonucleic acid (siRNA) molecule directing cleavage of severe acute respiratory syndrome (SARS) virus RNA through RNA interference, comprising a strand having nucleotide sequence with sufficient complementarity to the SARS virus RNA to direct cleavage of the SARS virus RNA through RNA interference, where the siRNA does not require the presence of nucleotides having a 2-hydroxy group for mediating RNA interference, and each strand of is 19-23 nucleotides in length. The siRNA is useful for modulating the expression of genes of associated with the development or maintenance of SARS virus infection, acute respiratory failure, viral pneumonia and/or disease states associated with SARS virus infection, and for treating SARS virus infection, acute respiratory failure, viral pneumonia and/or disease (transcription of SARS virus infection by preventing the strates associated with SARS virus infection by preventing the ctranscription of SARS sense. The siRNA is useful in diagnosis, and treatment of diseases and conditions the respond to the modulation of SARS sense. The siRNA is useful in ciagnosis, and therapeutic, diagnostic, target validation, genomic discovery, genetic capable of mediating RNA interference against SARS inside and ceconstituted in vitro system, and has high degree of specificity with respect to SARS RNA expression. The present sequence is a lower strand of constituted in vitro system, and has high degree of specificity with respect to SARS RNA expression. The present sequence is a lower strand of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                              Novel chemically synthesized double stranded short interfering nucleic acid molecule directing cleavage of severe acute respiratory syndrome virus RNA through RNA interference, useful for treating viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SARS; severe acute respiratory syndrome; siRNA; short interfering RNA; ss; RNA interference; gene silencing; SARS virus infection; acute respiratory failure; viral pneumonia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 0 A; 8 C; 1 G; 0 T; 10 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SARS coronavirus siRNA upper sequence 177.
                                                                                                                                                                                                                              Example 3; SEQ ID NO 1828; 219pp; English
                                                                                                          Haeberli P;
                                                                            SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    695 GAAGAAGCAGAGGAAGA 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADT63274 standard; RNA; 19 BP
23-MAY-2003; 2003US-00444853.
23-OCT-2003; 2003US-00693059.
24-NOV-2003; 2003US-00720448.
                                             14-JAN-2004; 2004US-00757803.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18 GAAGAAGAAGAGAAGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-APR-2004; 2004WO-US011320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-JAN-2005 (first entry)
                                                                                                         Mcswiggen J, Bharat C,
                                                                                                                                       WPI; 2004-766879/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SARS coronavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004092383-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-OCT-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADT63274;
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The invention relates to a chemically synthesised double stranded short interfering ribonucleic acid (siRNA) molecule directing cleavage of severe acute respiratory syndrome (SARS) virus RNA through RNA contressed comprising a strand having nucleotide sequence with sufficient complementarity to the SARS virus RNA to direct cleavage of the SARS virus RNA through RNA interference, where the siRNA does not require the presence of nucleotides having a 2'-hydroxy group for mediating RNA interference, and each strand of is 19-23 nucleotides in mediating RNA interference, and each strand of is 19-23 nucleotides in consociated with the development or maintenance of SARS virus infection, acute respiratory failure, viral pneumonia and/or disease states associated with SARS virus infection, and for treating SARS virus cute respiratory failure, viral pneumonia and/or disease states infection, acute respiratory failure, viral pneumonia and/or disease contents associated with SARS virus infection by preventing the transcription of SARS gene. The siRNA is useful in diagnosis, and transcription of SARS gene. The siRNA is useful in diagnosis, and transcription of SARS gene. The siRNA is useful in therefore, and pharmacogenomic applications. The siRNA is useful in therespeutic, diagnostic, target validation, genomic discovery, genetic capable of mediating RNA interference against SARS inside a cell or reconstituted in vitro system, and has high degree of specificity with respont to salve stranded siRNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                              Novel chemically synthesized double stranded short interfering nucleic acid molecule directing cleavage of severe acute respiratory syndrome virus RNA through RNA interference, useful for treating viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; TSA7005; Reg; pancreatic beta cell growth; hypoglycaemic; diagnosis; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.6%; Score 15.4; DB 1; Length 19; 94.1%; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 19 BP; 10 A; 1 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; SEQ ID NO 177; 219pp; English.
                                                                                                                                                                                                                                                          Haeberli P;
                                                                                                                                                                         (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 695 GAAGAAGCAGAGGAAGA 711
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23-OCT-2003; 2003US-00693059.
24-NOV-2003; 2003US-00720448.
14-JAN-2004; 2004US-00757803.
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Best Local Similarity 94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF82119 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16; Conservative
                                                                                                                                                                                                                                                              Bharat C,
                                                                                                                                                                                                                                                                                                                                           WPI; 2004-766879/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JP2001025389-A
                                                                                                                                                                                                                                                              Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF82119;
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18-JUL-1997;
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                                                                              11-MAY-1999
                                                                                              JP11032765-A
                                                                                                  09-FEB-1999
                                                                                          Synthetic.
                                                                          AAX18388;
                                                Query Match
                 TSA7005
                                                                  RESULT 1598
                                                                                      RT-PCR
                                                  Best Loca
Matches
                                                                    AAX18388,
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useful as primers for RT-PCR and determination of base seguences. The r
seguences allow for reproductive and highly efficient analysis of gene
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                                                           sednences
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                                                                                                                                                                Query Match
                                                                                                                                                                                            Local
                                                                                                                                                                                                                     Matches
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  ប្រម្ជាធិត្ត
ប្រម្ជាធិត្ត
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                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a PCR primer which is used in an example from the present invention for the isolation of human TSA7005 gene. The human TSA7005 protein shares 32% homology with human and mouse Reg proteins, and 34% homology with the rat Reg protein. TSA7005 has pancreatic beta cell growth activity and hypoglycaemic activity. The TSA7005 protein can be used for the diagnosis and treatment of diseases associated with the gene and its expression product
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                      TSA7005 gene, encoding a polypeptide useful for the diagnosis and treatment of diseases associated with its expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 15.2; DB 1; Length 16; 93.8%; Pred. No. 1.2e+03; Live 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 16 BP; 1 A; 0 C; 0 G; 14 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                               Example 1; Page 24; 25pp; Japanese.
                      99JP-00201279
                                                                      99JP-00201279
                                                                                                                           (SAKA ) OTSUKA PHARM CO LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 93,8
tes 15, Conservative
                                                                                                                                                                                 WPI; 2001-303742/32
                   15-JUL-1999;
                                                                      15-JUL-1999;
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DB 1; Length 17;

0.6%; Score 15.2;

Query Match

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The invention relates to a probe set for gene expression arrays to provide common equivalent signalling per probe and global amplification of the probe set has a pool of modified cDNA probes, each probe having a central target specific segment copied from a portion of a single mRNA transcript and a universal linker (a WRAP-Probe) located on or both terminal ends. The universal linker has reporter binding sites to join common reporters to the probes and primer binding sites to copy and amplify the probe. The probes and reporters are useful in componing discovery assays for a wide range of blomedical samples, including detection of nucleic acids and gene expression profiles in human diagnostics, forensics and genomic analysis. The methods are useful for amplifying and identifying any unknown DNA component and also for improving sensitivity with tissue microarrays or RNA arrays. The methods improve the quantification of gene expression and allow highly improved detection of rare transcripts or very small
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                                                                                                                                                                                                                                                                                                                                                                                                WRAP-Probe; gene expression array; global amplification; RNA array; ss;
tissue microarray; drug discovery assay; reporter binding site; forensic;
diagnostic; genomic analysis; universal linker; poly-T primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (WRAP
of
                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel probe sets with common universal linkers at one or both ends probes) for gene expression arrays to provide global amplification probe set and to provide common equivalent signaling regardless of
                                                                            ö
                                                                                                                                                                                                                                                                                                                                                           Modified Poly-T Primer #1 used in construction of probe sets.
                                       Length 17;
                                   0.6%; Score 15.2; DB 1; Length 1
93.8%; Pred. No. 1.2e+03;
ive 1; Mismatches 0; Indels
Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A; 0 C; 0 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 88; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENE-) GENETAG TECHNOLOGY INC.
                                                                                                             2708 TAAAAAAAAAAAA 2723
                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0187982P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-MAR-2001; 2001WO-US007508
                                                                                                                                           16 BAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                             AAS14174 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        construction of probe sets
                                                                          15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-596845/67.
                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200166802-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shafer DA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention comprises genes from Phanerochaete chrysosporium which exhibit a change in expression in the presence of an aromatic compound. The Phanerochaete chrysosporium genes of the invention are useful for analysing an environmental pollutant. The present DNA sequence represents an oligo dT PCR primer that was used in an example of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel genes of eukaryotic microorganism belonging to Phenerochate genus, and exhibiting change in expression of behavior in presence of aromatic compound, is useful for analyzing environmental pollutant.
                Gaps
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                                                                                                                                                                                                                                                                                            aromatic compound; gene expression alteration; environmental pollutant analysis; 8s; oligo dT; PCR; primer.
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0.6%; Score 15.2; DB 1; Length 19;
Best Local Similarity 93.8%; Pred. No. 1.3e+03;
Matches 15; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                        Environmental pollutant method-related oligo dT PCR primer.
Best Local Similarity 93.8%; Pred. No. 1.2e+03;
Matches 15; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 0 A; 0 C; 2 G; 15 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 9; 36pp; Japanese.
                                               ADM11779 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2002; 2002JP-00210632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUL-2002; 2002JP-00210632
                                                                                                                                                                                                                              20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-232127/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (WARI/) WARIISHI H.
(KUBI ) KUBOTA CORP.
                                                                                                                                                                                                                                                                                                                                                                           JP2004049103-A.
                                                                                                                                                                                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                            19-FEB-2004.
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ADM11779/c
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Gaps